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Regult
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Perfect score:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Database
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Maximum DB seq length: 200000000
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    OM protein - protein search, using sw model
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Total number of hits satisfying chosen parameters:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         on:
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                                                                                                                                                                                                                                                                                                                                                                                                                           Score
    Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Issued_Patents_AA:*

1: /ggn2_6/ptodata/1/iaa/5A_COMB.pep:*

2: /cgn2_6/ptodata/1/iaa/5B_COMB.pep:*

3: /cgn2_6/ptodata/1/iaa/6A_COMB.pep:*

4: /cgn2_6/ptodata/1/iaa/6B_COMB.pep:*

5: /cgn2_6/ptodata/1/iaa/PCTUS_COMB.pep:*

6: /cgn2_6/ptodata/1/iaa/backfiles1.pep:*
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Gapop 10.0 ,
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    Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries
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                                                                                                                                                                                                                                                                                                                                                                                                                           Length
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                 US-08-137-117D-86
US-08-436-717-86
PCT-US96-13152-3
US-09-157-370-3
US-08-887-352B-9
US-09-109-207C-10
US-09-109-20171-10
US-09-716-028-10
US-09-716-028-13
US-08-887-352B-13
US-08-867-352B-13
US-08-66-151-9
US-09-109-207C-13
US-09-109-207C-13
US-09-109-207C-13
US-09-109-207C-13
US-09-109-207C-13
US-09-109-207C-13
US-09-109-207C-13
US-09-109-207C-13
US-09-206-131-996-9
US-09-113-996-13
US-09-113-996-9
US-09-296-005-9
US-09-296-005-9
US-09-296-005-9
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US-09-113-996-9
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Sequence 86, Appl Sequence 2, Appli Sequence 3, Appli Sequence 10, Appli Sequence 11, Appli Sequence 13, Appli Sequence 13, Appli Sequence 13, Appli Sequence 13, Appli Sequence 14, Appli Sequence 15, Appli Sequence 16, Appli Sequence 17, Appli Sequence 18, Appli Sequence 18, Appli Sequence 19, Appli Sequence 19, Appli Sequence 19, Appli Sequence 10, Appli Sequence 2, Appli
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4. 5	44	43	42	41	40	39	ω 8	37	36	35	34	33	32	31	30	29	28
475	476	477	477	477	477	477	477	477.5	477.5	480.5	480.5	481	481	481	481	481	481
80.0	80.1	80.3	80.3	80.3	80.3	80.3	80.3	80.4	80.4	80.9	80.9	81.0	81.0	81.0	81.0	81.0	81.0
214	131	218	218	218	218	218	114	110	110	110	110	111	111	111	111	111	111
N	ω	4	4.	4.	w	w	N	4.	w	4.	w	4	4.	4.	w	ω	N
US-07-934-373C-39	US-08-579-378A-18	US-09-483-588-1	US-09L680-145-1	US-09-282-846-1	US-09-054-255-1	US-09-282-505-1	US-08-887-352B-8	US-09-025-403A-15	US-09-672-609-15	US-09-025-403A-17	US-09-672-609-17	US-10-113-996-7	US-09,-716-028-7	US-09-920-171-7	US-09-296-005-7	US-09-109-207C-7	US-08-887-352B-7
Sequence 39, Appl	Sequence 18, Appl	Sequence 1, Appli	Sequence 8, Appli	Sequence 15, Appl	Sequence 15, Appl	Sequence 17, Appl	Sequence 17, Appl	Sequence 7, Appli									

# RESULT 1 US-08-137-117D-86 US-08-137-117D-86 Sequence 86, Application US/08137117D Patent No. 5795965 GENERAL INFORMATION: APPLICANT: TSUCHIYA, Masayuki APPLICANT: BENDIG, Mary APPLICANT: BENDIG, Mary APPLICANT: JONES, Steven

ALIGNMENTS

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; TOPOLOGY: linear; MOLECULE TYPE: protein US-08-137-117D-86
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US-08-436-717-86
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US-08-436-717-86
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Best Local Similarity
Matches 100; Conserv
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Patent No. 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     GENERAL INFORMATION:
                                           INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 130 amino acid
TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                 PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/08/137,117
FILING DATE: 20-DEC-1993
APPLICATION NUMBER: WO PCT/JP92/00544
FILING DATE: 24-APR-1992
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTMARE: PatentIn Release #1.0,
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: SALDANHA, JOSE
TITLE OF INVENTION: RESHAPED HUMAN ANTIBODY
TITLE OF INVENTION: INTERLEUKIN-6 RECEPTOR
NUMBER OF SEQUENCES: 158
                                                                                                                                                            NAME: WEGNER, Harold C.
REGISTRATION NUMBER: 25,258
REFERENCE/DOCKET NUMBER: 534
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                  PRIOR APPLICATION DATA:
APPLICATION NUMBER: 19 3-95476
PILING DATE: 25-APR-1991
ATTORNEY/AGENT INFORMATION:
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PRIOR APPLICATION DATA:
APPLICATION NUMBER: JP 4-32084
FILING DATE: 19-FEB-1992
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                                 TOPOLOGY:
                                                                                                                                TELEFAX:
                                                                                                                                                TELEPHONE:
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CLASSIFICATION: 536
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               TYPE:
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3000 K Street, N.W
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JONES, Steven
SALDANHA, Jose
                                                               130 amino acids
                                                                                                                                (202) 672-5300
(202) 672-5399
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Pred. No. 2.7e-40;
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GENERAL INFORMATION:
                                                                                                                        Matches
                                                                                                                                                     Query Match
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                                                                                                                                                                                                                                                                             TELEFAX: (212) 838-3884
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                  PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08
FILING DATE: 27-Dec-95
APPLICATION NUMBER: EF
FILING DATE: 17-Aug-95
APPLICATION NUMBER: EF
                                                                                                                                                                                                                                                                                                                                                                        FILING DATE: 19-Sep-95
ATTORNEY/AGENT INFORMATION:
NAME: Norman D. Hanson
REGISTRATION NUMBER: 30,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                           REFERENCE/DOCKET NUMBER: BO
TELECOMMUNICATION INFORMATION:
TELEPHONE: (212) 688-9200
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER READABLE FORM: MEDIUM TYPE: 3.5" Co
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CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                       MOLECULE TYPE:
                                                                                                                                                                                                                 TYPE: amino acid
STRANDEDNESS: single
TOPOLOGY: linear
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           OPERATING SYSTEM:
SOFTWARE: ASCII
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    STREET: 805 Third
CITY: New York
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                                                                                                                                        Local
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                                                                                                                                                                                                                                                              ENGTH:
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61 GIPSRFSGSGSGTDFTLTISSLQPEDFATYYCQQSNEDPWTFGQGTKVEIKRTV 114
                            61 GVPSRFSGSGSGTDFTLTISSLQPEDFATYYCQHSWGIPRTFGQGTKVEIKRTV 114
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                                                                              1 DIQMTQSPSSLSASVGDRVTITTCRASQSVSTSSYSYMHWYQQKPGKAPKVLISYASNLES
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                                                                                                                                        Similarity
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                                                                                                                                                                                                                                                                   218
                                                                                                                          Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          IBM PC compatible SYSTEM: PC-DOS/MS-DOS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Attn: Norman D.
                                                                                                                                                                                                    protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    27-Dec-95
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90.1%;
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86.0%;
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Pred. No. 2.7e-40;
                                                                                                                                        Score 490; DB 5;
Pred. No. 2.9e-38;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Hanson
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APPLICANT: STEIPE, Boris
APPLICANT: STEIPE, Boris
ITITLE OF INVENTION: PROCESS FOR MODIFYING THE STABILITY OF ANTIBODIES
FILE REFERENCE: P8341-8072
CURRENT APPLICATION NUMBER: US/09/157,370A
CURRENT FILING DATE: 1998-09-21
EARLIER APPLICATION NUMBER: 08/765,179
EARLIER FILING DATE: 1997-01-14
EARLIER APPLICATION NUMBER: PCT/EP95/02626
EARLIER APPLICATION NUMBER: DCT/EP95/02626
EARLIER APPLICATION NUMBER: DCT/EP95/02626
EARLIER APPLICATION NUMBER: DCT/EP95/02626
EARLIER APPLICATION NUMBER: DCT/EP95/02626
EARLIER APPLICATION NUMBER: DC/P44 25 115.7
EARLIER FILING DATE: 1994-07-15
NUMBER OF SEQ ID NOS: 10
SOFTWARE: Patentin Ver. 2.1
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; TYPE: PRT
; ORGANISM: Homo sapiens
US-09-157-370-3
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Patent No. 5994511
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Best Local Similarity
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Patent No. 626223
TELEFAX: 650/952-9881
INFORMATION FOR SEQ ID NO: 9:
SEQUENCE CHARACTERISTICS:
LENGTH: 114 amino acids
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MEDIUM TYPE: 3.5 inch, 1.44 Mb floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: WinPatin (Genentech)
CURRENT APPLICATION DATA:
                                                                                                                                        APPLICATION NUMBER: US/08/887,352B FILING DATE: 03-Jul-1997 CLASSIFICATION: 530 ATTORNEY/AGENT INFORMATION: NAME: SVOBOda, Craig G. REGISTRATION NUMBER: 39,044 REFERENCE/DOCKET NUMBER: P1123
                                                                                          TELECOMMUNICATION INFORMATION: TELEPHONE: 650/225-1489
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CORRESPONDENCE ADDRESS:
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TITLE OF INVENTION:
TITLE OF INVENTION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Application US/08887352B
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RESULT 7
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US-08-887-352B-10
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Patent No. 5994511
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Best Local Similarity 86.0%;
Matches 98; Conservative
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Best Local Similarity
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             MEDIUM TYPE: 3.5 inch, 1.44 Mb floppy COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS SOFTWARE: WinPatin (Genentech) CURRENT APPLICATION DATA:
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NAME: Svoboda, Craig G.
REGISTRATION NUMBER: 39,0
REFERENCE/DOCKET NUMBER:
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TITLE OF INVENTION:
TITLE OF INVENTION:
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CORRESPONDENCE ADDRESS:
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TOPOLOGY: Linear
                                                                                                                                                                                                                                                                                             TYPE: Amino Acid
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Improved Anti-IgE Antibodies and Method
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Pred. No. 3.2e-38;
3; Mismatches 13
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Pred. No. 3.2e-38;
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US-09-109-207C-10
; Sequence 10, Application US/09109207C
; Patent No. 6172213

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RESULT 9
US-09-20-171-10
; Sequence 10, Application
; Patent No. 6682735
; GENERAL INFORMATION:
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; OTHER INFORMATION: Light chain sequence derived from MAB11
US-09-296-005-10
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US-09-296-005-10
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Henry B. Lowman, Leonard G. Presta, Paula M. Jardieu, John Lowe TITLE OF INVENTION: Improved Anti-IgE Antibodies and Method of Improving Polypeptide FILE REFERENCE: P1123R1 CURRENT APPLICATION NUMBER: US/09/109,207C CURRENT FILING DATE: 1998-06-30 PRIOR APPLICATION NUMBER: US 60/051,554 PRIOR FILING DATE: 1997-07-03 NUMBER OF SEQ ID NOS: 44

SEQ ID NO 10
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LENGTH: 114
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Best Local Similarity 86.0%;
Matches 98; Conservative
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Henry B. Lowman, Leonard G. Presta, Paula M. Jardieu, John Lowe TITLE OF INVENTION: Improved Anti-IgE Antibodies and Method of Improving Polypeptides FILE REPERENCE: PIL32CIr CURRENT APPLICATION NUMBER: US/09/296,005 CURRENT FILING DATE: 1999-04-21 EARLIER APPLICATION NUMBER: US/08/887,352 EARLIER FILING DATE: 1997-07-02 RARLIER FILING DATE: 1997-07-02 RARLIER FILING DATE: 1997-07-02
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ORGANISM: Artificial FEATURE: NAME/KEY: Artificial
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TYPE: PRT
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Pred. No. 3.2e-38;
3; Mismatches 13
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APPLICANT: Presta, Leonard G.
APPLICANT: Joardieu, Paula M.
APPLICANT: Jowe, John
TITLE OF INVENTION: Improved Anti-IgE Antibodies (as ame
FILE REFERENCE: P1113C2US
CURRENT APPLICATION UNBER: US/09/920,171
CURRENT FILING DATE: 2001-08-01
PRIOR APPLICATION NUMBER: US 08/887,352
PRIOR APPLICATION NUMBER: US 09/296,005
PRIOR APPLICATION NUMBER: US 09/296,005
PRIOR FILING DATE: 1999-04-21
PRIOR FILING DATE: 1999-04-21
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US-09-716-028-10
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APPLICANT: Henry B. Lowman, Leonard G. Presta, Paula M. Jardieu, John Lowe
APPLICANT: Henry B. Lowman, Leonard G. Presta, Paula M. Jardieu, John Lowe
TITLE OF INVENTION: Improved Anti-IgE Antibodies and Method of Improving Polypeptide
FILE REFERENCE: P1123R1

CURRENT APPLICATION NUMBER: US/09/716,028

CURRENT FILING DATE: 2000-11-17

PRIOR APPLICATION NUMBER: US 09/109,207

PRIOR FILING DATE: 1998-06-30

PRIOR APPLICATION NUMBER: US 60/051,554

PRIOR APPLICATION NUMBER: US 60/051,554

PRIOR FILING DATE: 1997-07-03

NUMBER OF SEQ ID NOS: 44

SEQ ID NO 10

LENGTH: 114
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SEQ ID NO 10
LENGTH: 114
TYPE: PRT
ORGANISM: Artificial Sequence
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Patent No. 6723833
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                      TYPE: PRT
ORGANISM: Artificial
FEATURE:
                                                                                                                                                                                                                                                                                            NAME/KEY: Artificial LOCATION: 1-114
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61
                        61 GVPSRFSGSGSGTDFTLTISSLQPEDFATYYCQHSWGIPRTFGQGTKVEIKRTV 114
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                                                                                                           1 DIQMTQSPSSLSASVGDRVTITCRASQSVSTSSYSYMHWYQQKPGKAPKVLISYASNLES
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GVPSRFSGSGSGTDFTLTISSLQPEDFATYYCQQSHEDPYTFGQGTKVEIXRTV 114
                                                                                   DIQLTQSPSSLSASVGDRVTITCRASQSVDYDGDSYMNWYQQXPGKAPKLLIYAASYLES
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Pred. No. 3.2e-38;
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Pred. No.
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RESULT 11

US-10-113-996-10

Application US/10113996

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RESULT 12
US-08-887-352B-13
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US-10-113-996-10
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PRIOR FILING DATE: 2001-08-01
NUMBER OF SEQ ID NOS: 44
SEQ ID NO 10
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Sequence 10, App.
No. 6761889
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FILE REFERENCE: P1123C3US
CURRENT APPLICATION NUMBER: US/10/113,996
CURRENT FILING DATE: 2002-04-01
PRIOR APPLICATION NUMBER: US 08/887,352
PRIOR PILING DATE: 1997-07-02
PRIOR PILING DATE: 1997-07-02
PRIOR FILING DATE: 1999-04-21
PRIOR FILING DATE: 1999-04-21
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             CLASSIPICATION: 530
ATTORNEY/AGENT INFORMATION:
NAME: Svoboda, Craig G.
REGISTRATION NUMBER: 39,044
REFERENCE/DOCKET NUMBER: P112
TELECOMMUNICATION INFORMATION:
TELEPHONE: 650/252-1489
TELEPHAY: 650/952-9881
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APPLICANT: Henry
INFORMATION FOR SEQ
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ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                     COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5 inch, 1.44 Mb floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                       SOFTWARE: WinPatin (Genentech)
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/887,
FILING DATE: 03-Jul-1997
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TITLE OF INVENTION:
NUMBER OF SEQUENCES:
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Local Similarity 86.0%;
nes 98; Conservative
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Lowe, John
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Pred. No. 3.2e-38;
3; Mismatches 13
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Patent No. 6037453
GENERAL INFORMATION:
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Best Local Similarity 86.0%;
Matches 98; Conservative
                                                                        TELEFAX: 650/952-9881 INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                               FILING DATE: 06-Jun-19
APPLICATION NUMBER: 08
FILING DATE: 15-MAR-19
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08
FILING DATE: 26-JAN-19
                                                                                                     NAME: Svoboda, Craig G. REGISTRATION NUMBER: 39,044. REFERENCE/DOCKET NUMBER: POTELECOMMUNICATION INFORMATION: TELEPHONE: 650/225-1489
                                                                                                                                                                                               FILING DATE: 07-MAY-1992
PRIOR APPLICATION DATA:
APPLICATION UNMBER: 07/70
FILING DATE: 14-AUG-1991
ATTORNEY/AGENT INFORMATION:
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CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/466,
FILING DATE:
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MEDIUM TYPE: 3.5 inch, 1.44 Mb
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                     SEQUENCE CHARACTERISTICS
                                                                                                                                                                                                                                                                                                                                                                                                                                           CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/466163
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APPLICANT: Presta, Leonard
TITLE OF INVENTION: Immunog
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LENGTH: 218 amino acids
TYPE: Amino Acid
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APPLICATION NUMBER:
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TOPOLOGY:
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                 ENGTH: 218 amino acids
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Pred. No. 6.8e-38;
3; Mismatches 13
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Query Match

81.8%;

Score 486;

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Length 218;

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GRGANISM: Artificial
FEATURE:
FEATURE:
FACTURE:

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US-09-109-207C-13
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                                                                                                                                                                                                                                                                                                                                                         Sequence 13, Application US/09296005
Patent No. 6290957
Patent No. 6290957
Patent No. 6290957
Patent No. 6290957

GENERAL INFORMATION:
APPLICANT: Henry B. Lowman, Leonard G. Presta, Paula M. Jardieu, John Lowe
TITLE OF INVENTION: Improved Anti-IgE Antibodies and Method of Improving Polypeptides
FILE REFERENCE: P1123C1r
CURRENT APPLICATION NUMBER: US/09/296,005
CURRENT FILING DATE: 1999-04-21
EARLIER APPLICATION NUMBER: US 08/887,352
EARLIER FILING DATE: 1997-07-02
NUMBER OF SEQ ID NOS: 26
SEQ ID NO 13
LENGTH: 218
TYPE: NEW
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APPLICANT: Henry B. Lowman, Leonard G. Presta, Paula M. Jardieu, John Lowe
APPLICANT: Henry B. Lowman, Leonard G. Presta, Paula M. Jardieu, John Lowe
TITLE OF INVENTION: Improved Anti-IgE Antibodies and Method of Improving Polypeptide
FILE REFERENCE: P1123R1
CURRENT APPLICATION NUMBER: US/09/109,207C
CURRENT FILING DATE: 1998-06-30
PRIOR APPLICATION NUMBER: US 60/051,554
PRIOR PPLICATION NUMBER: US 60/051,554
PRIOR FILING DATE: 1997-07-03
NUMBER OF SEQ ID NOS: 44
SEQ ID NO 13
LENGTH: 218
TYPE: PRT
ORGANISM: Artificial
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Matches 98; Conserv
   Query Match
Best Local Similarity
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LOCATION: 1-218
COTHER INFORMATION: Light chain sequence derived from MAB11
                                                                                                                                                                                                                                                                                                                                          TYPE: PRT
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Score 486; DB 3;
Pred. No. 6.8e-38;
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Search completed: December 29, 2004, 19:37:36 Job time : 28.3991 secs

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1: geneseqp1980s:*
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3: geneseqp2001s:*
4: geneseqp2001s:*
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348.266 Million cell updates/sec
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

## SUMMARIES

Result

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Result No.	Score	Query Match	Length	80	ID	Description
1	594		114	σį	AAE28148	Aae28148 Humanised
ผ	501	84.3	111	N	AAR85242	N
w	494	•	218	IJ	AAM49204	
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10	488	•	238	ω	AAW90932	Aaw90932 Humanised
11	488	•	238	υį	ABB74901	•
12	488		240	4	AAB45995	Aab45995 Human MUC
13	486	•	114	N	AAW13922	Aaw13922 Light cha
14	486	•	114	N	AAW95654	Aaw95654 Mus muscu
15	486	•	114	N	AAW95655	Aaw95655 Mus muscu
16	486	•	114	4.	AAB76944	Aab76944 Variable
17	486		114	æ	ADN07031	
18	486		218	N	AAW95658	_
19	486		218	w	AAY85200	Aay85200 Light cha
20	486		218	4.	AAB76947	_
21	486		218	œ	ADN07034	Anti
22	486		240	4	AAB45996	
23	485		238	ω	AAW90931	Aaw90931 Humanised
24	485		238	v	ABB74900	Abb74900 Humanised
25	484		233	8	ADL93655	Adl93655 Human CD4

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81.1	81.1	81.3	81.3	81.3	81.3	81.3	81.3	81.3	81.3	81.3	81.3	81.3	81.3	81.3	81.3	81.3	81.3	81.3	81.5
108	108	240	240	240	240	240	240	240	240	240	240	219	114	114	108	108	108	108	240
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Ad192461	Abp96010	Ad192369	Abj38594	Abp95997	Aab45994	Aab46005	Aab46006	Aab46008	Aab46038	Aab46007	Aay02472	Adn07066	Adn07030	Aab76943	Ado36372	Ad192386	Ad191345	Abp96009	Aab46010
Single do	HSA antib	Human pha	Hepatitis	Human ser	Human MUC	Human MUC	Human MUC	Human MUC	Human TF	Human MUC			Anti-IgE	Variable	! Intracell	Anti-HSA	VL chain	HSA antib	Human MUC

# ALIGNMENTS

# RESULT 1 AAE28148

AAE28148 standard; protein; 114 ₽

AAE28148;

27-DEC-2002 (first entry)

Humanised 9F3 version 13 (V13) light chain variable domain.

Human; interferon alpha; IFNalpha; insulin-dependent diabetes mellitus; autoimmune disease; systemic lupus erythematosus; autoimmune thyroiditis; antibody therapy; immunosuppressive; antiinflammatory; dermatological; 9F3 monoclonal antibody; light chain variable domain.

Homo sapiens.

Synthetic

Key Region Region Region Location/Qualifiers 24. .38 /note= "Complementarity 93. .102 54. .60 /note= "Complementarity determining region 1 determining region 2 (CDR1) " (CDR2) "

WO200266649-A2

note=

"Complementarity determining region 3 (CDR3)"

29-AUG-2002.

29-JAN-2002; 2002WO-US002709.

22-FEB-2001; 2001US-0270775P. 09-JAN-2002; 2002US-00044896.

(GETH ) GENENTECH INC.

Chuntarapai A, Kim JK, Presta LG, Stewart H

WPI; 2002-682767/73.

New anti-interferon alpha monoclonal antibody, 9F3, useful for treating autoimmune diseases e.g. insulin-dependent diabetes mellitus, systemic lupus erythematosus and autoimmune thyroiditis.

Disclosure; Page 97; 100pp; English.

The present invention relates to novel anti-interferon alpha (IFNalpha)

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Matches
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Best Local
Antibodies (Ab) comprising a humanised variable region specifically binding to a varicella zoster virus (VZV) can be used to treat or prevent VZV infection. They may optionally be attached to a therapeutic agent. They may also be used, when labelled, to detect VZV particles and infected cells in blood; for the removal/ neutralisation of infectious VZV in blood; for the selection/ isolation of human monoclonal Abs and for the design of (non-) peptide mimetics with similar diagnostic and therapeutic uses. The variable region CDR portion is derived from murine Ab 206 and the heavy and light chain variable region frameworks are NEWM
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             monoclonal antibody, 9F3 which binds to and neutralises a biological activity of IFN-alpha subtypes. The anti-IFNalpha monoclonal antibodies of the invention are useful for treating autoimmune diseases such as insulin-dependent diabetes mellitus, systemic lupus erythematosus and autoimmune thyroiditis. They are useful as reagents in diagnostic assays for IFN-alpha expression, for the affinity purification of various IFN-alpha expression, for the affinity purification of various for IFN-alpha expression of the combinant cell culture or natural resources and for the detection of IFN-alpha in diagnostic assay methods. Sequences of the invention are also useful in antibody therapy. The present sequence is anti-luman IFNalpha monoclonal antibody 9F3 version 13 (V13) light chain
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Antibody; varicella zoster virus; Ab; VSV; treatment; prevention; infection; detection; isolation; monoclonal antibody; MAb; mimetic; humanised; murine; heavy chain; light chain; variable region; CDR;
                                                                                                                                                                                           Claim 7; Page
                                                                                                                                                                                                                                              Varicella zoster virus gpIII antibody with humanised variable region for treatment, prevention or diagnosis of varicella zoster virus
                                                                                                                                                                                                                                                                                                           WPI; 1996-010932/01
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             GVPSRFSGSGSGTDFTLTISSLQPEDFATYYCQHSWGIPRTFGQGTKVBIKRTV
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     GVPSRFSGSGSGTDFTLTISSLQPEDFATYYCQHSWGIPRTFGQGTKVEIKRTV
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        DIQMTQSPSSLSASVGDRVTITCRASQSVSTSSYSYMHWYQQKPGKAPKVLISYASNLES
                                                                                                                                                                                                                                                                                                                                              Wallace TP,
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                                                                                                                                                                                         31; 58pp; English
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                                                                                                                                                                                                                                                                                                                                                                                                                        94US-00234133
                                                                                                                                                                                                                                                                                                                                                                                   BIOPHARMACEUTICALS
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Pred. No. 2.9e-32;
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                                                                                                                                   or prevent
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RESULT 3
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                                                                                                                                                                                                                                                                                                                                                     Monoclonal antibody; mAb; humanised; murine; mouse; 5c8; hu5c8; light chain; anti-CD145; CD145-antibody complex; 3D structure; three dimensional structure; drug design; drug discovery; activated T cell; CD40 interaction; T cell dependent immune response; agonist; antagonist; immune response; inflammatory response; autoimmune disease; allergy; inhibitor response; organ graft rejection; B cell cancer; Alzheimer's disease; multiple sclerosis; antiinflammatory; immunosuppressive; antiallergic; cytostatic; dermatological; antiasthmatic; nootropic; neuroprotective; antiarteriosclerotic; antiriviral; antidabetic; cardiant; antiischaemic; vasodilator; antirheumatic; antiarthritic; antipsoriatic; immunomodulator; antibody; complementarity determining region; CDR; protein co-ordinate data.
                                                                                                                                                                                                                                                                        Key
Region
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28-JUN-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            and REI respectively. The framework may include at least one mutation that improves binding specificity or affinity. The heavy and light che variable regions are preferably (VZVHu-VH, -VHKFIIS, -VHTLY, -VHKVRSE -VHS) and VZVHuVK respectively
                                                                                                                                                                                       Binding-
                                                                                                                                                                                                                                                 Region
                                                                                                                                                                                                                                                                                                             Chimeri
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAM49204 standard;
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                                 07-MAR-2002.
                                                                                          Binding-site
                                                                                                                             Region
                                                                                                                                                                 Region
                                                                                                                                                                                                             Binding-site
                                                                                                                                                                                                                                                                                                                        Homo
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                                                       WO200218445-A2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         98;
                                                                                                                                                                                        -site
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                                                                                                                                                                                                                /note=
                                                                                          95.
                                                                                                                                                                                                                                                                                   Location/Qualifiers
                                                                                                     /label= CDR3
/note= "Complementarity
                                                                                                                                       /label= CDR2
/note= "Complementarity
                                                                                                                                                                                                    note=
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                                                                                                                                                                                                                                      label= CDR1
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                                                                                "Binds
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Pred. No. 4e-2
4; Mismatches
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                                                                                                                                                                                                                                                              part of the crystal of the invention"
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                                                                                CD145
                                                                                                                                                                            CD145 (AAM49202) "
                                                                                                                                                                                                    CD145
                                                                                (AAM49202)
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RSE or
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31-AUG-2001; 2001WO-US027352

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RESULT 4
AAW22842
ID AAW2
XX
AC AAW2
XX
AC 12-S
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                                                                                                                                                                                                                                                                                                                                                                                                                                       CC complex with an anti-CD45 antibody or its antigen-hinding fragment, and CC comprises human CD145 (AAM49202) and a humanised version of the murine CC comprises human CD145 (AAM49202) and a humanised version of the murine CC comprises human CD145 (AAM49202) and a humanised version of the murine CC comprises human CD145 (AAM49202) and a humanised version of the murine CC control (pp39) F-BAM, Sc 8 antigen, CD40CR and TRAP) is 32 kD type II CC macrophages with CD40 which is expressed on activated T CC cells. It interacts with CD40 which is expressed on activated T CC cells. This CD40:CD145 interaction is required for T cell dependent CC antibody responses, type I T-helper cell response and nature B cells, CC cells. This CD40:CD145 interaction is required for T cell dependent CC artivities of macrophages, and disruption of the CD40:CD145 interaction coll derivated for T cell dependent CC artivities of macrophages, and disruption of the CD40:CD145 interaction coll derivated for T cell dependent CC artivities of an anti-CD145 antibody has been shown to reduce the symptoms of autoimmune and inflammatory conditions. The crystal structure of the invention can be used to determine the three dimensional structure CC for the CD145 anti-CD145 antibody complex, and thereby provide information cc about this interaction which may be of use in designing non-antibody CC colls agonists and antagonists which modulate the CD40:CD145 interaction. CC Such compounds may be used in the treatment of an unwanted immune cc asthma, chronic obstructive pulmonary response, an autoimmune disease, an allergy, an inhibitor response to a thraspeutic agent, rejection of a CC cashama, chronic obstructive pulmonary should be compounded to compounds may be used in the treatment of an unwanted immune cc asthma, chronic obstructive pulmonary should be compounded to compound the collection of a collection of 
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Best Local S
Matches 91
    12-SEP-1997
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                                           AAW22842;
                                                                                 AAW22842 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 2002-329760/36
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         and Alzheimer's disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Кагривав М,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (BIOJ ) BIOGEN INC.
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                                                                                                                                                                                                                                                                                                                                                        91,
                                                                                                                                                                                                                                                                                                                                                                           Similarity
                                                                                                                                                                                                            GVPSRFSGSGSGTDFTLTISSLQPEDFATYYCQHSWGIPRTFGQGTKVEIKRTV
                                                                                                                                                                                      GVPARFSGSGSGTDFTLTISSVEPEDFATYYCQHSWEIPPTFGGGTKLEIKRTV
                                                                                                                                                                                                                                                                    DIVLTQSPATLSVSPGERATISCRASQRVSSSTYSYMHWYQQKPGQPPKLLIKYASNLES
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                                                                                                                                                                                                                                                                                                                                                                                                                                       218 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                5c8 (hu5c8).
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                                                                                                                                                                                                                                                                                                                                                      Conservative
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2001US-0276452P.
  (first entry)
                                                                                 protein; 132
                                                                                                                                                                                                                                                                                                                                                                       83.2%;
79.8%;
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                                                                                                                                                                                                                                                                                                                                                      13;
                                                                                                                                                                                                                                                                                                                                                                         Score 494; DB 5
Pred. No. 2e-25;
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Best Local
L-selectin; humanised antibody; HuDreg 55; acute organ damage; organ failure; poly-trauma; haemorrhagic-traumatic shock.
                                                                                                                                                                                                                                                                                                                                                           (MAb) light chain variable region, useful in medicine, pharmacology and biochemistry. The isotype of a MAb secreted by the human/human hybridoma HT was determined to be mu and kappa. Human MAb was purified, and the antigen recognised by human MAb CLN"-IgM identified by western blotting
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human; tumour antigen; cancer; monoclonal; antibody; livariable region; medicine; pharmacology; biochemistry; complementarity determining region.
                                     Humanised
                                                            17-OCT-2003
03-JUN-1997
                                                                                                  AAW13563;
                                                                                                                         AAW13563 standard;
                                                                                                                                                                                                                                                                                                                                     Sequence 132 AA;
                                                                                                                                                                                                                                                                                                                                                                                                  The present sequence is a human anti-tumour antigen monoclonal (MAb) light chain variable region, useful in medicine, pharmaco
                                                                                                                                                                                                                                                                                                                                                                                                                                      Claim 9; Page 11; 14pp; Japanese
                                                                                                                                                                                                                                                                                                                                                                                                                                                               Anticancer human monoclonal antibody variable region sequences related DNA and RNA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                03-OCT-1995;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human anti-tumour antigen antibody light chain variable region
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                                                                                                                                                                                                    61 GVPSRFSGSGSGTDFTLTISSLQPEDFATYYCQHSWGIPRTFGQGTKVEIKRTV 114
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                                                                                                                                                                                                                                                                                      97;
                                                                                                                                                                                                                                                                                                   Similarity
                                                                                                                                                                                                                                                   DIQMTQSPSSLSASVGDRYTITCRASQSVSTSSYSYMHWYQQKPGKAPKVLISYASNLES
                                                                                                                                                                                                                                      DIQMTQSPSSLSASVGDRVTITCRASQSIS----SYLNWYQQKPGKAPKLLIYAASSLQS
                                                                                                                                                                                     GVPSRFSGSGSGTDFTLTISSLQPEDFATYYCQQSYSTPQTFGQGTKVEIKRTV
                                    anti-L-selectin antibody HuDreg 55 light chain.
                                                                                                                                                                                                                                                                                      Conservative
                                                            (revised)
(first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 95JP-00278266
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/label=_CDR_2
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                                                                                                                        protein;
                                                                                                                                                                                                                                                                                                83.0%;
85.1%;
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                                                                                                                         218
                                                                                                                                                                                                                                                                                    Score 493; DB
Pred. No. 1.6e
8; Mismatches
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                                                                                                                                                                                                                                                                                   DB 2; L
l.6e-25;
les 5;
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iochemistry; CDR;
                                                                                                                                                                                                                                                                                                            Length 132;
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RESULT 6
ABP90 XX
ABP90 XX
ABP90 XX
ABP9 XX
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             17-AUG-1995;
19-SEP-1995;
27-DEC-1995;
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Homo sapiens.
Chimeric.
                                                                                                                     Ligand; human serum albumin; HSA; antibody; cytostatic; anti-HIV; antiinflammatory; antianaemic; immunosupproseive; neuroprotective; dual-specific ligand; cancer; HIV infection; hepatitis; rubella; inflammation; autoimmune disorder; multiple sclerosis; Crohn's dis
                                                Homo sapiens
                                                                                              myasthenia gravis.
                                                                                                                                                                                                                                                                                                  01-MAY-2003
                                                                                                                                                                                                                                                                                                                                                                                                ABP96011 standard;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        anti-selectin antibody to prevent acute organ damage and failure - during extracorporeal circulation or following rauma, e. g. haemorrhagic-traumatic shock.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          61
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BOEHRINGER
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                                                                                                                                                                                                                                            related Vkappa
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95EP-00114696.
95US-00578953.
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Pred. No. 3.7e-25;
4; Mismatches 12
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                                                                                                                     a; anaemia;
disease;
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RESULT 7
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(C) (a) a first single immunoglobulin variable domain with a binding of specificity to a first antigen or epitope; and (b) a second complementary immunoglobulin single variable domain with a binding activity to a second the first and second domain with a binding activity to a second of the first and second domains are mutually complementary, and the first and second domains lack mutually complementary domains that share the same specificity. (I) has cytostatic, anti-HIV, antianaemic, anti-HIV, ant
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
Best Local 9
                  immunoglobulin single variable domain; cytostatic; antiinflammatory; antirheumatic; antiarthritic; antiathmatic; antiallergic; antibacterial; virucide; immunosuppressive; antidiabetic; neuroprotective; muscular; dermatological; gene therapy; inflammatory; rheumatoid arthritis; asthma; Crohn's disease; allergic hypersensitivity; bacterial; viral infection; autoimmune disorder; type I diabetes; multiple sclerosis; myasthenia gravis; systemic lupus erythematosus; cancer; light chain variable region; human; E5sd.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Dual-specific ligand having immunoglobulins with binding specificity to different antigens or epitopes, useful for treating, preventing or diagnosing diseases, e.g. cancer, HIV infection, inflammations, or
                                                                                                                                                                                                                                                       Single domain antibody-related light chain variable region
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                                                                                                                                                                                                                                                                                                                                                                                                             standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    GVPSRFSGSGSGTDFTLTISSLQPEDFATYYCQHSWGIPRTFGQGTKVEIKR
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85.7%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Matches
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Best Local
nephrotropic; antiinfertility; neuroprotective; antiarteriosclerotic; hepatotropic; humanized; apoptosis; systemic lupus erythematosus; Hashimoto disease; rheumatoid arthritis; graft versus host disease; Sjorgen's syndrome; anemia; Addison's disease; scleroderma; sterility; Goodpasture syndrome; Crohn's disease; sterility; myasthenia gravis;
                                                                              Pas; antibody; human; anti-inflammatory; anti-anemic; antidiabetic; anti-allergic; anti-arthritic; antiviral; immunomodulatory; cardiant; dermatological; immunosuppressive; thyromimetic; antitheumatic; anti-Fas;
                                                                                                                                                      Humanised anti-Fas designed light chain Leu 1 protein.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Example 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      New dual-specific ligands, useful in drug discovery and development, for diagnosing, preventing or treating a disease, such as cancer, autoimmune disease, or inflammatory disease, including rheumatoid
                                                                                                                                                                                                                                                         AAW90930 standard; protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          arthritis or asthma.
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27-DEC-2002;
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                                                                                                                                                                                                                                                                                                                                                                                                                                  Fig 3; 174pp; English
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Pred. No. 2.4e-25;
5; Mismatches 6
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                                                                                                                                                                                                                        This invention describes a novel humanized anti-Pas antibody-like CC molecule (1) that, induces apoptosis in cells with an abnormal Fas/Fas CC ligand system, by binding to Fas on the cell surface, and prevents CC apoptosis in cells with a normal system, by inhibiting binding between CC Pas and its ligand. The products of the invention have anti-inflammatory, CC anti-anemic, antidiabetic, anti-allergic, anti-arthritic, antiviral, CC antirheumatic, nephrotropic, anti-infertility, neuroprotective, CC antirheumatic, nephrotropic, antiinfertility, neuroprotective, CC antirheumatic, nephrotropic, antiinfertility, neuroprotective, CC antisted side to cell surface Fas or inhibit it by competitive CC diseases associated with the Fas/Fas ligand system, especially systemic CC lugus erythematosus, Hashimoto disease, rheumatoid arthritis, graft CC versus host disease, Sjorgen's syndrome, pernicious or hypoplastic CC anemia, Addison's disease, scleroderma, Goodpasture syndrome, Crohn's CC disease, autoimmune hemolytic anemia, sterility, myasthenia gravis, CC disease, autoimmune hemolytic anemia, sterility, myasthenia gravis, CC dependent diabetes mellitus, allergy, arteriosclerosis, myocarditis, CC cardiomyopathy, glomerulomephritis, hepatitis (filminant, chronic, viral (B, C or D) or alcoholic), and transplant rejection. (I) selectively cells. They bind to both human and murine Fas, so can be evaluated in murine disease models. (I) act on the active site of Fas, i.e. they mimic channed human set anti-Fas antibody light chain construct designated Leu 1 cc which is described in the method of the invention
                                                                                                                                               Matches
                                                                                                                                                                Query Match
Best Local
                                                                                                                                                                                                                      Sequence 238
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             New humanized anti-Fas antibody, useful for inflammatory or autoimmune disease, induces cells with abnormal Fas-Fas ligand systems.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           multiple sclerosis; Basedow's disease; thrombopenia purpura; allergy; insulin dependent diabetes mellitus; arteriosclerosis; myocarditis; cardiomyopathy; glomerulonephritis; hepatitis; transplant rejection.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Claim
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          30-SEP-1998;
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81
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                                                                                                                                               98;
                                                                                                                                                                Similarity
                                                                                         DIQMTQSPSSLSASVGDRVTITCRASQSVSTSSYSYMHWYQQKPGKAPKVLISYASNLES
GVPSRFSGSGSGTDFTLTISSLQPEDFATYYCQQSNEDPRTFGQGTKVEIKRTV
                                GVPSRFSGSGSGTDFTLTISSLQPEDFATYYCQHSWGIPRTFGQGTKVEIKRTV 114
                                                                      DIVLTQSPSSLSASVGDRVTITCKASQSVDYDGDSYMWYQQKPGKAPKLLIYAASNLBS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Page 156-157; 263pp;
                                                                                                                                               Conservative
                                                                                                                                                                                                                      ¥,
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98JP-00276882.
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                                                                                                                                                              82.3%;
86.0%;
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                                                                                                                                               4
                                                                                                                                           Score 489; DB 3;
Pred. No. 4.7e-25;
4; Mismatches 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  treating or preventing e.g. apoptosis selectively in
                                                                                                                                               12;
                                                                                                                                                                                Length 238
                                                                                                                                               Indels
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 134
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RESULT 10
AAW90932
ID AAW90
XX
AC AAW90
XY
DT 08-AU
DT 08-AU
XX
DE Humar
XX
KW Fas;
KW earti-
KW derm,
KW neph,
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The invention relates to a preventive or treating agent for diseases caused by abnormality in Fas/Fas ligand system containing as the active component an antibody having as the light chain subunit a polypeptide containing residues 1-218 of one of 3, 239 residue amino acid sequences, or residues 1-451 of one of 3, 470 residue amino acid sequences, all fully defined in the specification and having an activity of combining specifically with mammalian Fas and an activity of inducing apoptosis in a cell expressing Fas. The agent has immunosuppressive and antiallergic activity and is used for preventing and treating autoimmune diseases,
            Fas; antibody; human; anti-inflammatory; anti-anemic; antidiabetic; anti-allergic; anti-rathritic; antiviral; immunomodulatory; cardiant; dermatological; immunosuppressive; thyromimetic; antirheumatic; anti-Fas; nephrotropic; antiinfertility; neuroprotective; antiarteriosclerotic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human; mouse; humanised anti-Fas antibody; Fas/Fas light chain subunit; apoptosis; immunosuppressive; autoimmune disease; allergy; atopic.
                                                                                                                    08-AUG-2000
                                                                                                                                                                            AAW90932 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    29-MAR-2000; 2000JP-00090918
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   28-MAR-2001; 2001JP-00093106
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           JP2001342148-A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Humanised anti-Fas antibody light chain SEQ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           26-APR-2002
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  containing humanized anti-Pas antibody, used ting autoimmune diseases, allergy, and atopy.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             2002-145113/19
                                                                                                                                                                                                                                                    81
                                                                                                                                                                                                                                                                               61
                                                                                                                                                                                                                                                                                                              21
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                                                                                                                                                                                                                                                                                                                                                                     98;
                                                                                                                                                                                                                                                                                                                                                                                    Similarity
                                                                                                                                                                                                                                                                                                                             DIQMTQSPSSLSASVGDRVTITCRASQSVSTSSYSYMHWYQQXPGKAPKVLISYASNLES
                                                                                                                                                                                                                                                                                                                                                                                                                                                          atopy and
                                                                                                                                                                                                                                                                              GVPSRFSGSGSGTDFTLTISSLQPEDFATYYCQHSWGIPRTFGQGTKVEIKRTV 114
                                                                                                                                                                                                                                                                                                                                                                                                                               238 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   standard;
                                                                                                                                                                                                                                                    GVPSRFSGSGSGTDFTLTI
                                                                                                                                                                                                                                                                                                            DIVLTQSPSS
                                                                                                                                                                                                                                                                                                                                                                     Conservative
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                                                                                                                  (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              humanised anti-Fas antibody; Fas/Fas
                                                                                                                                                                           protein;
                                                                                     designed light chain Leu 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                            others
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                                                                                                                                                                                                                                                                                                                                                                                   82.3%;
apoptosis;
                                                                                                                                                                            238
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                                                                                                                                                                                                                                                    SSLOPEDFATYYCOOSNEDPRIFGOGTKVEIKRTV
                                                                                                                                                                                                                                                                                                                                                                                   Score 489; DB 5;
Pred. No. 4.7e-25;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                allergy,
                                                                                                                                                                                                                                                                                                                                                                   Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              IJ
                                                                                                                                                                                                                                                                                                                                                                   12;
lupus erythematosus;
                                                                                     protein.
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antiallergic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               preventing
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GIPSRFSGSGSGTDFTLTISSLQPEDFATYYCQQSNEDPRTFGQGTKVEIKRTV

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                                                                                                                                                                                                                   CC molecule (I) that, induces apoptosis in cells with an abnormal Fas/Fas CC ligand system, by binding to Fas on the cell surface, and prevents CC apoptosis in cells with a normal system, by inhibiting binding between CC Fas and its ligand. The products of the invention have anti-inflammatory, CC anti-anemic, antidiabetic, anti-allergic, anti-arthritic, anti-viral, cimunomodulatory, dermatological, immunosuppressive, thyromimetic, CC antiarteriosclerotic, cardiant and hepatropic activity. (I) induce approves by binding to cell surface Fas or inhibit it by competitive CC inhibition of ligand binding. (I) are used to treat and/or prevent CC diseases associated with the Fas/Fas ligand system, especially systemic CC upus erythematosus, Hashimoto disease, rheumatoid arthritis, graft CC versus host disease, soleroderma, germicious or hypoplastic CC anemia, Addison's disease, soleroderma, Goodpasture syndrome, Crohn's CC disease, autoimmune hemolytic anemia, sterility, mysathenia gravis, CC multiple sclerosis, Basedow's disease, thrombopenia purpura, insulin CC dependent diabetes mellitus, allargy, arteriosclerosis, myocarditis, CC cardiomyopathy, glomerulonephritis, hepatitis (fulminant, chronic, viral CC (B, C or D) or alcoholic), and transplant rejection. (I) selectively inhibit apoptosis in normal calls but selectively induce it in abnormal CC cells. They bind to both human and murine Fas, so can be evaluated in murine disease models. (I) act on the active site of Fas, i.e. they mimic CC inducing a human anti-murine antibody response. This sequence represents a humanised anti-Fas antibody light chain construct designated Leu 3 cc which is described in the method of the invention
                                                                                                                                Best Loc
Matches
                                                                                                                                                                        Query Match
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                                                                                                                                                                                                                Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           New humanized anti-Fas antibody, useful for inflammatory or autoimmune disease, induces
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Hashimoto disease; rheumatoid arthritis; graft versus host disease;
Sjorgen's syndrome; anemia; Addison's disease; scleroderma; sterili
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Claim 3; Page 161-162; 263pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Serizawa N,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               30-SEP-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        inflammatory or autoimmune disease,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (SANY ) SANKYO CO LTD.
                                                                                                                                                     Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           2000-258930/23.
                                                  21
                                                                                                                                Similarity
97; Conserv
GVPSRFSGSGSGTDFTLTISSLQPEDFATYYCQHSWGIPRTFGQGTKVEIKRTV 114
                                                  DIVLTQSPSSLSASVGDRVTITCKASQSVDYDGDSYMNWYQQKPGKAPKLLIYAASNLES
                                                                       DIQMTQSPSSLSASVGDRVTITCRASQSVSTSSYSYMHWYQQKPGKAPKVLISYASNLES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     abnormal Fas-Fas ligand systems.
                                                                                                                                  Conservative
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98JP-00276882
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                                                                                                                                                   82.2%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         a novel humanized anti-Fas antibody-like
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                                                                                                                                Score 488; DB
Pred. No. 5.4e
5; Mismatches
                                                                                                                                                   488; DB 3;
No. 5.4e-25;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        treating or preventing apoptosis selectively i
                                                                                                                                                                    Length 238;
                                                                                                                                  Indels
                                                                                                                                0,
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in
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RESULT 12
AAB45995
ID AAB45
XX
XX
AC AAB45
AC AAB45
AC AAB45
XX
XX
DT 23-MA
XX
DB Human
XX
XW MUCl;
KW antii
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ABB74901
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                                                                                                                                                                                                                                                                                                                                Matches
                                                                                                                                                                                                                                                                                                                                              Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                              The invention relates to a preventive or treating agent for diseases caused by abnormality in Fas/Fas ligand system containing as the active component an antibody having as the light chain subunit a polypeptide containing residues 1-218 of one of 3, 239 residue amino acid sequences, or residues 1-451 of one of 3, 470 residue amino acid sequences, all fully defined in the specification and having an activity of combining specifically with mammalian Fas and an activity of inducing appotesis in a cell expressing Fas. The agent has immunosuppressive and antiallergic activity and is used for preventing and treating autoimmune diseases,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 2002-145113/19.
N-PSDB; ABL45941.
MUC1; human; vaccine; conformation-dependent antigen; antibody; cancer; antidiotypic antibody; cytostatic; virucidal; antibacterial; TF antige
                                            Human MUC-1
                                                                                                                                   AAB45995 standard; peptide; 240 AA
                                                                                                                                                                                                                                                                                                                                                                                         Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Example 14 (Preparatory); Page 94; 194pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Drug containing humanized anti-Fas antibody, used for preventing
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      29-MAR-2000; 2000JP-00090918
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       light chain subunit; apoptosis; immu autoimmune disease; allergy; atopic.
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                                                                          23-MAR-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (SANY ) SANKYO
                                                                                                                                                                                                                                        61
                                                                                                                                                                                                                                                                      21
                                                                                                                                                                                                             81
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     mouse; humanised anti-Fas antibody; Fas/Fas ligand;
chain subunit; apoptosis; immunosuppressive; antiallergic;
                                                                                                                                                                                                                                                                                               1 DIQMTQSPSSLSASVGDRVTITCRASQSVSTSSYSYMHWYQQKPGKAPKVLISYASNLES 60
                                                                                                                                                                                                                                                                                                                                            Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                autoimmune diseases, allergy, and atopy.
                                                                                                                                                                                                         GIPSRFSGSGSGTDFTLTISSLQPEDFATYYCQQSNEDPRTFGQGTKVEIKRTV 134
                                                                                                                                                                                                                                        GVPSRFSGSGSGTDFTLTISSLQPEDFATYYCQHSWGIPRTFGQGTKVEIKRTV
                                                                                                                                                                                                                                                                   238 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                     atopy and
                                            scFv clone R2.
                                                                                                                                                                                                                                                                                                                              Conservative
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                                                                          (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                      others
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85.1%;
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                                                                                                                                                                                                                                                                                                                                           Score 488; DB 5;
Pred. No. 5.4e-25;
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                                                                                                                                                                                                                                                                                                                              12;
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                                                                                                                                                                                                                                                                                                                                                          Length 238;
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                                                                                                                                                                                                                                        114
                                                                                                                                                                                                                                                                                                                              Gaps
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RESULT 13 AAW13922

8X5

AAW13922

AAW13922 standard; protein; 114 AA

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                                                                                                                                                                                                                                                                                            CC of an antidiotypic antibody (AD2) or another peptide which: (a) encoder of specifically binds to the binding size of an antibody (AD1) or an antigen CC binding molecule; and (b) immunologically mimics the initial antigen. The CP compete is partially or completely conformation-dependent, and has an CC acids. (I) is used in the form of linear or circular naked DNA and/or CC with a viral vector and/or adjuvants. The products of the invention have CC cytostatic, virucidal, antibacterial and antiparasitic. The invention have CC are not proteins or peptides, as defined above but which have epitopes CC which show an immunogenic structure; (2) preparing (V1) and (V2); (3) CC which show an immunogenic structure; (2) preparing (V1) and (V2); (3) CC which show an immunogenic structure; (2) preparing (V1) and (V2); (3) CC which show an immunogenic structure; (2) preparing (V1) and (V2); (3) CC which show an immunogenic structure; (2) preparing (V1) and (V2); (3) CC which show an induction of structure; (3) preparing (V1) and (V2); (3) CC which show an induction of structure; (4) MUC1-conformation epitope CC all fully defined in the specification; (4) MUC1-conformation epitope CC mimics having one of 31 approximately 60 residue amino acids sequences, all fully in the specification; (6) Treatbohydrate epitope mimetics for the specification; and (7) DNA sequences and fully defined in the specification; (6) Treatbohydrate epitope mimetics cancer, and infectious diseases, e.g. caused by prions, viruses, bacteria and parasites. The vaccines are effective in cases where vaccination has cc previously not been possible
                                                                                                                                                                        Matches
                                                                                                                                                                                                                 Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         This invention describes a novel vaccine (VI) against conformation-dependent antigens (CDA) comprising DNA (I) and/or an antibody, or peptide which immunologically imitates CDA, is new. (I) encodes a region of an antiidiotypic antibody (Ab2) or another peptide which: (a) specifically binds to the binding site of an antibody (Ab1) or an antiger
                                                                                                                                                                                                                                                            Sequence 240
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Disclosure; Page 5-9; 36pp; German.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Vaccines against conformation-dependent or non-peptide antigens, based on DNA encoding peptide which mimics the antigen, useful e.g. as antitumor \mathbf{p}
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 2001-049937/06
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              27-MAY-1999;
09-SEP-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      antiparasitic; infectious disease
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189
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                                                                                                                                                                        96;
                                                                                                                                                                                               Similarity
                                        GVPSRFSGSGSGTDFTLTISSLQPEDFATYYCQHSWGIPRTFGQGTKVEIKR 112
                                                                                    DIQMTQSPSSLSASVGDRVTITCRASQSIS----SYLNWYQQKPGKAPKLLIYHASFLQS
                                                                                                          DIQMTQSPSSLSASVGDRVTITCRASQSVSTSSYSYMHWYQQKPGKAPKVLISYASNLES 60
GVPSRFSGSGSGTDFTLTISSLQPEDFATYYCQQPWEPPRTFGQGTKVEIKR
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99DE-01043016
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                                                                                                                                                                        Score 488; DB 4;
Pred. No. 5.5e-25;
6; Mismatches 6
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Best Local (
                                                                                                                                Variable light chain; IgE; antibody; anti-IgE; reduction; preve histamine; production; hypersensitivity; allergen; anaphylaxis; atopic allergy; asthma; allergic rhinitis; conjunctivitis; hay
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              invention preferably contains the sequence represented by AAW13912 in the complementarity determining region-1 (CDR-1) of the heavy chain variable region. The antibody of the invention also contains the sequence represented by AAW13913 in the CDR-3 of the light chain variable region. The antibody is capable of binding to adr type hepatitis B (HB) virus antigen. A human anti-HB virus monoclonal antibody preparation which is highly safe and is effective to adr type HB virus can be provided, using
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Antibody; heavy chain; light chain; variable region; human; monoclonal; complementarity determining region; human; adr type hepatitis B virus; HB virus; CDR; virus antigen; anti-HB antibody; vaccine.
     WO9901556-A2
                                                          Mus musculus
                                                                                                              eczema; anaphylactic shock; urticaria.
                                                                                                                                                                                                                                                        Mus musculus anti-IgE e426
                                                                                                                                                                                                                                                                                                                  08-JUN-1999
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Claim 7;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              anti-Hepatitis B antibody - used in a
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              VPSRPSGSGSGTDFTLTISSLQPEDFATYYCQQSYSTPRTFGQGTKVEIKRTV
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          VPSRFSGSGSGTDFTLTISSLQPEDFATYYCQHSWGIPRTFGQGTKVEIKRTV 114
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             LEMTQSPSSLSASVGDRVTITCRASQSIS----SYLWWYQQKPGKAPKLLIYAASSLQSG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Page 10-11;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     IQMTQSPSSLSASVGDRVTITCRASQSVSTSSYSYMHWYQQKPGKAPKVLISYASNLESG
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                                                                                                                                                                                                                                                     variable light chain.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 486; DB 2
Pred. No. 4e-25;
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                                                                                                                                                                                             anti-IgE; reduction; prevention;
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RESULT 15
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                                                                                                                                                                                                     Variable light chain; IgE; antibody; anti-IgE; reduction; prevention; histamine; production; hypersensitivity; allergen; anaphylaxis; atopic allergy; asthma; allergic rhinitis; conjunctivitis; hay fever;
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            Lowman HB,
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                                                                                                                                                                    Mus musculus
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                                     (GETH ) GENENTECH INC.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Improving affinity of polypeptides, particularly anti-IgE antibodies - by identifying aspartyl residues which undergo isomerisation and substituting alternative residues and screening for affinity against the
                                                                                                                                                                                                                                                                                                          Sequence 114 AA;
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l Similarity 86.0%;
98; Conservative
                                                                                            GVPSRFSGSGSGTDFTLTISSLQPEDFATYYCQHSWGIPRTFGQGTKVBIKRTV 114
                                                                         GVPSRFSGSGSGTDFTLTISSLQPEDFATYYCQQSHEDPYTFGQGTKVEIKRTV 114
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Ig kappa chai Ig kappa chai
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Query Match Local

Length 123;

C;Accession R;Klein, R. Bur. J. Imm A;Title: Exx A;Title: Exx A;Reference A;Accession A;Status: p A;Molecule A;Residues: A;Cross-refi C;Superfami C;Keywords: F;32-106/Do	RESULT S40331 Ig kap C;Spec C;Date		45	443	42	41	40	ى 8 س	37	36	35	34	33	32	31	30
C; Accession: S40331 Eur. J. Immunol. 23, 3248-3271, 1993 A; Title: Expressed human immunoglobu A; Reference number: S40312; MUID:940 A; Accession: S40331 A; Status: preliminary; translation n A; Status: preliminary; translation n A; Molecule type: mRNA A; Mesidues: 1-123 < KLE> A; Cross-references: EMBL:X72441; NID C; Superfamily: immunoglobulin V regi C; Kuywords: heterotetramer; immunoglobulin homo	RESULT 1 S40331 Ig kappa chain - human C;Species: Homo sapiens C;Date: 06-Mar-1994 #sec		432.5	4 4 3 3 3 3	433	434	434	436	437	437	437	437	437.5	438	438	438
; S40331 ; Jaenichen, l j Jaenichen,	human sapiens 1994 #seq		72.8	72.9 72.9		73.1		73.4		73.6	73.6	73.6	73.7	73.7	73.7	73.7
R.; Za 18-3271 1 immun 112; MU 1:ransla 1:X7244 bbulin imer; im	(man) uence_			126				111								108
H.G. lin chi genes 80891; PMID:8; 80891; PMID:8; ot shown ot shown :g441350; PID on; immunoglol obulin logy <imm></imm>	man Lens (man) #seguence_revision 26-May-1995 #t	ALIGNMENTS		2 S40335 2 S40369				2 S52789 1 KVMS84			1 K1HUBN	1 K1HUAU			2 \$40350	1 К1НИКА
nes and their hypermutation. D:8258341 PIDN:CAA51109.1; PID:g441351 globulin homology	#text_change 21-Jan-2000		kappa	Ig kappa chain - h	kappa chain	kappa chain -	kappa chain V	Ig kappa chain V r	kappa chain V	kappa chain	kappa chain		_	kappa	kappa	Ig kappa chain V-I

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RESULT 2
$40367
Ig kappa chain V-J-C region - human
C;Species: Homo sapiens (man)
C;Date: 19-May-1994 #sequence_revision 26-May-1995 #text_change 21-Jan-2000
C;Accession: $40367
R;Klein, R.; Jaenichen, R.; Zachau, H.G.
Bur: J. Immunol. 23, 3248-3271, 1993
A;Title: Expressed human immunoglobulin chi genes and their hypermutation.
A;Reference number: $40312; MUID:94080891; PMID:8258341
A;Accession: $40367
                                      C;Superfamily: immunoglobulin V region; immunoglobulin homology C;Keywords: heterotetramer; immunoglobulin F;33-107/Domain: immunoglobulin homology <IMM>
                                                                                                                    A;Status: preliminary; translation
A;Molecule type: mRNA
A;Residues: 1-127 <KLE>
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95; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                    GVPSRFSGSGSGTDFTLTISSLQPEDFATYYCQQSYSTPRTFGQGTKVEIK 123
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Pred. No. 1.1e-32;
7; Mismatches 5
  Score 482;
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R;Klein, R.; Jaenichen, K.; Zachar, T...
Eur. J. Immunol. 23, 3248-3271, 193
A;Title: Expressed human immunoglobulin chi genes and their hypermutation.
A;Reference number: $40312; MUID:94080891; PMID:8258341
A;Accession: $40370
A;Accession: $40370
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S40370
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                A;Experimental source: thymic B lymphocytes
A;Note: sequence extracted from NCBI backbone (NCBIN:113208, NCBIP:113209)
C;Superfamily: immunoglobulin V region; immunoglobulin homology
F;16-90/Domain: immunoglobulin homology < TMM's
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Eur. J. Immunol. 22, 2231-2236, 1992
A;Title: Human monoclomal striational autoantibodies isolated from thymic B lymphocytes A;Reference number: A49047; MUID:92387224; PMID:1516616
A;Accession: B49047
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Ig kappa chain V region (monoclonal striational autoantibody StrAB SA-1A) - G;Species: Homo sapiens (man) C;Date: 19-Dec:1993 #sequence_revision 18-Nov-1994 #text_change 09-Jul-2004 C;Accession: B49047
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A; Residues: 1-108 < VIC>
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Ig kappa chain -
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;Date: 06-Mar-1994 #sequence_revision 26-May-1995 #text_change
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93; Conserv
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                                                                                 DIOMTOSPSSLSASVGDRVTITCRASOSVSTSSYSYMHWYQQXPGKAPKVLISYASNLES
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GVPSRFSGSGSGTDFTLTISSLQPEDFATYYCQHSWGIPRTFGQGTKVEIKR 112
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                                       DIQLTQSPSSLSASVGDRVTITCRASQSIST----FLHWYQQNLGKAPKLLIYAASNLQS
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83.0%;
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Pred. No. 7e-32;
7; Mismatches
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Pred. No. 5.2e-32;
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R;Portolano, S.; Chazenbalk, G.D.; Hutchison, S.J.; McLachlan, S.M.; Rapoport, submitted to the EMBL Data Library, June 1992
A;Description: Lack of promiscuity in autoantigen-specific H and L chain combiners. S31977
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                A;Map position: 2p12-2p12
C;Complex: an immunoglobulin heterotetramer subunit consists of two identical light hain disulfide bonds; in some cases, such as IgA and IgM, the subunits associate int C;Superfamily: immunoglobulin V region; immunoglobulin homology C;Keywords: heterotetramer; immunoglobulin c;Keywords: heterotetramer; immunoglobulin p;16-90/Domain: immunoglobulin homology cIMM>
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FEBS Lett. 222, 6-10, 1987
A;Title: Chemical modification of the carboxyl groups of protein substrates enhances th. A;Reference number: S02572; MUID:88005152; PMID:3115831
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R;Watanabe, S.; Hilschmann, N.
HOppe-Seyler's Z. Physiol. Chem. 351, 1291-1295, 1970
A;Title: The primary structure of a monoclonal kappa-type
A;Reference number: A01868; MUID:71032830; PMID:4097974
A;Accession: A01868
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C;Date: 24-Apr-1984 #sequence_revision 02-Jul-1998 #text_change 09-Jul-2004
C;Accession: A01868; S02574
S
                                                                                                                                                                                 A;Cross-references: EMBL:Z15081; NID:g38501; PIDN:CAA78790.1; PID:g38502
C;Superfamily: immunoglobulin V region; immunoglobulin homology
C;Keywords: heterotetramer; immunoglobulin
                                                                                                                                                                                                                                                             A; Molecule type: mRNA
A; Residues: 1-109 < POR>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             C;Species: Homo sapiens (man)
C;Date: 06-Feb-1995 #sequence_revision 06-Feb-1995 #text_change 21-Jan-2000
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A;Note: the C region of this chain has the
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                                                                                                                                                        F;16-90/Domain: immunoglobulin homology <IMM>
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       A;Gene: GDB:IGKV1
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A; Residues: 1-108 <WAT>
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Best Local
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Best Local (
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       61 GVPSRFSGSGSGTDFTLTISSLQPEDFATYYCQHSWGIPRTFGQGTKVEIKR 112
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  1 DIQMTQSPSSLSASVGDRVTITCRASQSVSTSSYSYMHWYQQKPGKAPKVLISYASNLES 60
                                                                               Similarity
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82.1%;
                                                 77.3%; Score 459; DB 2; 79.6%; Pred. No. 8.7e-31; tive 11; Mismatches 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (Hau) -
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Pred. No. 4.9e-31;
8; Mismatches 8
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                                                                                                       Length 109
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Eur. J. Immunol. 23, 3248-3371, 1993
A;Title: Expressed human immunoglobulin chi genes and their hypermutation.
A;Reference number: $40312; MUID:94080891; PMID:8258341
A;Accession: $40352
A;Status: preliminary; translation not shown
A;Molecule type: mRNA
A;Residues: 1-131 <KLE>
A;Cross-references: RMNA:
A;Cross-references: RMNA: Vregion; immunoglobulin homology
C;Keywords: heterotetramer; immunoglobulin
F;36-110/Domain: immunoglobulin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Ig kappa chain V-J-C region - human
C;Species: Homo sapiens (man)
C;Date: 19-May-1994 #sequence_revision 26-May-1995 #text_change
C;Accession: S40352
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   RESULT 8
S40352
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Ig kappa chain V region - human
()Species: Homo sapiens (man)
()Date: 13.7an.1995 #sequence_revision 13-Jan.1995 #text_change 24-May-2001
()Accession: $44122
R; Hawkins, R. E.; Zhu, D.; Ovecka, M.; Winter, G.; Hamblin, T.J.; Stevenson, submitted to the EMBL Data Library, March 1994
A;Description: Idiotypic vaccination against human B-cell lymphoma: rescue of A;Reference number: $44105
A;Recession: $44122
A;Status: preliminary
A;Mocession: $44122
A;Status: preliminary
A;Mocession: $44122
A;Status: preliminary
A;Mocession: $44126
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A;Cross-references: EMBL:Z31390; NID:g472976; PIDN:CAA83265.1;
C;Superfamily: immunoglobulin V region; immunoglobulin homology
C;Keywords: heterotetramer; immunoglobulin
F;16-90/Domain: immunoglobulin homology <IMM>
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                                                           GVPSRFSGSGSGTDFSLTISSLQPEDVATYYCQKYNSVPRTFGQGTKVEIKRTV
                                                                                            GVPSRFSGSGSGTDFTLTISSLQPEDPATTYCQHSWGIPRTFGQGTKVEIKRTV 114
                                                                                                                                         DIQMTQSPSSLSASVGNRVTITCRASQGIS----NYLAWYQQKPGKVPKLLIYAASTLQS
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82.1%; Pred. No. 1e-30;
vative 7; Mismatches
                                                                                                                                                                                                                                        76.9%;
79.8%;
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                                                                                                                                                                                                                                        Score 457; DB 2;
Pred. No. 1.5e-30;
                                                                                                                                                                                                                         Mismatches
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                                                                                                                                                                                                                                                           Length 131
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Ig kappa chain - human
(¿Species: Homo sapiens (man)
(¿Species: Homo sapiens (man)
(¿Species: Homo sapiens (man)
(¿Appa char-1994 #sequence_revision 26-May-1995 #text_change
(¿Accession: $40317
R;Klein, R.; Jaenichen, R.; Zachau, H.G.
Eur. J. Immunol. 23, 3248-3271, 1993
Eur. J. Immunol. 23, 3248-3271, 1993
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R;Reininger, L.; Shibata, T.; Ozaki, S.; Shirai, T.; Jaton, J.C.; Izui, R;Reininger, L.; Shibata, T.; Ozaki, S.; Shirai, T.; Jaton, J.C.; Izui, Bur. J. Immunol. 20, 771-777, 1990
A;Title: Variable region sequences of pathogenic anti-mouse red blood conference number: S09955; MUID:90269328; PMID:2347362
A;Recession: S09963
A;Recession: S09963
A;Residues: 1-111 | REI | REI | RESIDENCE | RESIDE
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A;Accession: S31981
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A;Reference number: S40312; MUID:94080891; I
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C;Species: Mus musculus (house mouse)
C;Datc: 12-Feb-1993 #sequence_revision 12-Feb-1993 #text_change 09-Jul-2004
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C;Superfamily: immunoglobulin V region; immunoglobulin homology
C;Keywords: heterotetramer; immunoglobulin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 R;Portolano, S.; Chazenbalk, G.D.; Hutchison, submitted to the EMBL Data Library, June 1992
                                                                                                                                                                                                                                                                                                      RESULT 11
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ;Superfamily: immunoglobulin V region; immun;Keywords: heterotetramer; immunoglobulin;16-94/Domain: immunoglobulin homology <IMM>
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89
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83; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                      GVPSRFSGSGSGTDFTLTISSLQPEDFATYYCQHSWGIPRTFGQGTKVEIK 111
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Pred. No. 1.8e-30;
9; Mismatches 11
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      genes and their
PMID:8258341
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Ig kappa chain C; Species: Homo C; Date: 06-Jan-: C; Accession: S4'
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S40333
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                                                  A;Status: preliminary; translation not shown
A;Molecule type: mRNA
A;Molecule type: mRNA
A;Residues: 1-125 <KLE>
A;Cross references: EMBL:X72443; NID:g441354; PIDN:CAA51111.1; PID:g441355
A;Cross references: EMBL:X72443; NID:g441354; PIDN:CAA51111.1; PID:g441355
C;Superfamily: immunoglobulin V region; immunoglobulin homology
C;Keywords: heterotetramer; immunoglobulin
                                                                                                                                                                                     EUR. J. Immunol. 23, 3248-3271, 1993
A;Title: Expressed human immunoglobulin chi
A;Reference number: S40312; MUID:94080891; E
A;Accession: S40333
                                                                                                                                                                                                                                                                         C;Accession: S47182
C;Accession: S47182
C;Accession: S47182
C;Accession: S47182
C;Accession: A.P.
Submitted to the EMBL Data Library, June 1994
Submitted to the EMBL Data Library, June 1994
A;Description: Cloning and analysis of IgM anti-thyroglobulin autoantibodies from patien
A;Reference number: S47181
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              A;Cross-references: EMBL:X72427; NID:g441322; PIDN:CAA51095.1; PID:g441323 C;Superfamily: immunoglobulin V region; immunoglobulin homology
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A;Molecule type: mRNA
A;Residues: 1-129 <KLE>
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;Date: 06-Jan-1995 #sequence_revision 06-Jan-1995 #text_change
;Accession: S47182
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                                              ;34-108/Domain:
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Best Local Similarity
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37-111/Domain: immunoglobulin homology <IMM>
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80.2%;
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Pred. No. 3.9e-30;
2; Mismatches 8
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Pred. No. 3.1e-30;
8; Mismatches 10
  Score 448;
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PMID:8258341
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GVPSRFSGSGSGTDFTLTISSLQPEDFATYYCQHSWGIPRTFGQGTKVEIK

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A;Molecule type: mRNA
A;Residues: 1-106 <ESP>
A;Crose-references: EMBL:X75387
A;Crose-references: EMBL:X75387
C;Superfamily: immunoglobulin V region; immunoglobulin
F;16-90/Domain: immunoglobulin homology <IMM>
F;24-34/Region: complementarity determining region 1
F;50-56/Region: complementarity determining region 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gene 148, 167-168, 1994
A;Title: Phage display of a human antibody
A;Reference number: PC2397; MUID:95011651;
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                                                                                                                                                           A;Cross-references: EMBL:X72424
C;Superfamily: immunoglobulin V region; immunoglobulin
C;Keywords: heterotetramer; immunoglobulin
                                                                                                                                                                                                                                                                                    R;Klein, R.; Jaenichen, R.; Zachau, H.G.
Eur. J. Immunol. 23, 3248-3271, 193
A;Tille: Expressed human immunoglobulin chi genes
A;Reference number: S40312; MUID:94080891; PMID:82
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C;Date: 13-Jun-1995 #sequence_revision 14-Jul-1995
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                                                                                                                                         C;Keywords: heterotetramer; immunoglobuli F;31-105/Domain: immunoglobulin homology
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C;Date: 06-Mar-1994 #sequence_revision 26-May-1995
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                                 DIOMTOSPSSLSASVGDRVTITCRASOSVSTSSYSYMHWYQQKPGKAPKVLISYASNLES
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                                                                                         Score 447; DB 2;
Pred. No. 9.2e-30;
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Pred. No. 7.3e-30;
8; Mismatches 8;
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9; Mismatches
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8; Mismatches
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PMID:8258341
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Search completed: December 29, 2004, 19:36:34 Job time : 24.9742 secs

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KV1E HUMAN
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ALIGNMENTS	AAH17870	Q6PJB5	KV3T_MOUSE	KV1L_HUMAN	KV3U_MOUSE	KV1R_HUMAN	AAH34146	Q6PIH4	KV1C_HUMAN	KV1D HUMAN	KV1T_HUMAN	Q9UL79	KV1M_HUMAN	KV1F HUMAN
	Aah17870 homo sapi	Q6pjb5 homo sapien	P01672 mus musculu	P01604 homo sapien	P01673 mus musculu	P01610 homo sapien	Aah34146 homo sapi	Q6pih4 homo sapien	P01595 homo sapien	P01596 homo sapien	P01612 homo sapien	Q9ul79 homo sapien	P01605 homo sapien	P01598 homo sapien

#### Query Match Best Local Similarity Matches 92; Conserva Clin. Immunol. Immunopathol. 87:184-192(1998). EMBL; AF035037; AAD56273.1; -. PIR; B49047; B49047. PIR; S34083; S34083. HSSP; P01607; 1BWW. InterPro; IPR007;110; Ig-11ke. InterPro; IPR003596; Ig-v. Pfam; PF00047; 1g; 1. SMART; SM00406; IGv; 1. PROSITE; PS50835; IG\_LIKE; 1. Q9UL77; Q9UL77; 01-MAY-2000 01-MAY-2000 SEQUENCE FROM N.A. MEDLINE-99277139; PubMed=9614934; Mu X., Liu B., Van der Merwe P.L., Young D.C.; Homo sapiens (Human). Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo. MCBI\_TaxID=9606; 01.MAY-2000 (TrEMBLrel. 13, Created) 01.MAY-2000 (TrEMBLrel. 13, Last sequence update) 01-CT-2003 (TrEMBLrel. 25, Last annotation update) Myosin-reactive immunoglobulin light chain variable "Myosin-reactive autoantibodies in rheumatic carditis and normal fetus."; 61 GVPSRFSGSGSGTDFTLTISSLQPEDFATYYCQHSWGIPRTFGQGTKVEIKR 1 DIQMTQSPSSLSASVGDRVTITCRASQSVSTSSYSYMHWYQQKPGKAPKVLISYASNLES GVPSRFSGSGSGTDFTLTISSLQPEDFATYYCQQSYSTSWTFGEGTKVEIKR 108 DIQMTQSPSSLSASVGDRVTITCRASQSIS----SYLNWYQQKPGKAPNLLIYAASSLQS 108 108 AA; 78.1%; Score 464; DB 2; Length 108; larity 82.1%; Pred. No. 1.7e-37; Conservative 8; Mismatches 8; Indels PRELIMINARY; 108 11738 MW; C06681716C4D16F3 CRC64; PRT; Kalis N.N., Berney S.M., 108 region 4. 56

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MEDLINE=22388257; PubMed=12477932;

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Strausberg R.L., Feingold E.A., Grouse L.H., Derge J.G., Schuler G.D.,

Klausner R.D., Collins F.S., Wagner L., Schaefer C.F., Bhat N.K.,

Altschul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Hsieh F.,

Altschul S.F., Zordan H., Moore T., Max S.I., Wang J., Hsieh F.,

Hopkins R.F., Jordan H., Moore A.A., Rubin G.M., Hong L.,

Niatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,

Niatchenko L., Marusina K., Bonaldo M.F., Casavant T.L., Scheetz T.E.,
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05-JUL-2004
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Homo sapiens (Human).
Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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InterPro; IPR003596; Ig_v.
Pfam; PF00047; ig; 1.
SMART; SM00406; IGv; 1.
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PROSITE; PS50835; IG_LIKE;
DD-structure; Bence-Jones;
Immunoglobulin V region.
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GO:0005576; C:extracellular; NF
GO:0003823; F:antigen binding;
GO:0006955; P:immune response;
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response; NAS.
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Last sequence Last anno
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Pred. No. 2.7e
8; Mismatches
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                                                                                                                                                                                                                                                                                                                                                        sequence update) annotation updat
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NAS.
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                                                                                                                                                                                                                                                                                                                                                        update)
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PIR; PIR;

B49047; PH0867; S16840;

PH0867. S16840.

rheumatic carditis human anti-antibody V region genes.", J. Immunol. 161:2020-2031(1998)

SEQUENCE FROM N.A.

MEDLINE=98375893; PubMed=9712075;

Adderson E.E., Shikhman A.R., Ward K.B., Cunningham M.W.;

Molecular analysis of polyreactive monoclonal antibodies from

anti-N-acetylglucosamine/anti-myosin

NCBI\_TaxID=9606;

Hominidae;

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RESULT
Q96SA9
ACCOME DIT
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Best Local (
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Q96SA9;
01-DEC-2001
01-DEC-2001
01-MAR-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        InterPro; IPRO03599; Ig.
InterPro; IPRO07110; Ig-like.
InterPro; IPRO03597; Ig.cl.
InterPro; IPR003006; Ig_MHC.
InterPro; IPR003006; Ig_V.
Pfam; PP070549; Cl. set; 1.
Pfam; PP070547; Ig; 2.
SMART; SM00407; IG; 2.
SMART; SM00406; IGV; 1.
SMART; SM00406; IGV; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Brownstein M.J., Usdin T.B., Toshiyuki S., Carninci P., Prange Raha S.S., Loquellano N.A., Peters G.J., Abramson R.D., Mullah Bosak S.A., McEwan P.J., McKernan K.J., Malek J.A., Gunaratne Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Huly Villalon D.K., Muzny D.M., Sodergren E.J., Lu X., Gibbs R.A., Fahey J., Helton E., Ketteman M., Madan A., Rodrigues S., Sanc Whiting M., Madan A., Young A.C., Shevchenko Y., Bouffard G.G. Blakesley R.W., Touchman J.W., Green E.D., Dickson M.C., Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M., Butterfie Krzywinski M.I., Skalska U., Smailus D.B., Schnerch A., Schein
                                                                                                                                                           Eukaryota; Metazoa;
Mammalia; Eutheria;
                                                                                                                                                                                                          01-DEC-2001 (TrEMBLrel. 19, Created)
01-DEC-2001 (TrEMBLrel. 19, Last sequence update)
01-MAR-2004 (TrEMBLrel. 26, Last annotation updat
Anti-streptococcal/anti-myosin immunoglobulin kap
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PROSITE; PS50835; IG_LIKE; PROSITE; PS00290; IG_MHC; UHypothetical protein.
SEQUENCE 236 AA; 25707 N
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Strausberg R.;
Submitted (JUN-2004) to the
EMBL; BC073764; AAH73764.1;
                                                                                                                                                                                                 variable region (Fragment)
                                                                                                                                                                                  Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SEQUENCE FROM N.A.
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                                                                                                                                                                                                                                                                                                                                        GVPSRFSGSGSGTDFTLTISSLQPEDFATYYCQQAHSFPFTFGPGTKVDIKRTV
                                                                                                                                                                                                                                                                                                                                                                                        DIQMTQSPSSVSASVGDRVTITCRASQGIS----SWLAWYQQKPGKAPKLLIYAASSLQS
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                                                                                                                                                                                   (Human)
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Primates;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 459;
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                                                                                                                                                            Craniata; Vo
Catarrhini;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              99:16899-16903 (2002)
                                                                                                                                                                                                        st annotation update) immunoglobulin kappa
                                                                                                                                                                                                                                                                                                                                                                                                                                          Mismatches
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naratne P.
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G.G.,
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Y S.J.,
P.H.,
k S.W.,
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RESULT
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REPLINE=22388257; PubMed=12477932;

REPLINE=22388257; PubMed=12477932;

REPLINE=22388257; PubMed=12477932;

RA Klausner R.D., Collins F.S., Wagner L., Shenmen C.M., Schuler G.D.,

RA Altechul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,

RA Altechul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,

RA Altechul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,

RA Altechul S.F., Jordan H., Moore T., Max S.I., Mang J., Hsieh F.,

RA Hopkins R.F., Jordan H., Moore T., Max S.I., Mang L., Hong L.,

RA Hopkins R.F., Jordan H., Moore T., Casavant T.L., Scheetz T.E.,

RA Bounstein M.J., Usdin T.B., Bonaldo M.F., Casavant T.L., Scheetz T.E.,

RA Brownstein M.J., Usdin T.B., Toshiyuki S., Carninci P., Prange C.,

RA Raha S.S., Loquellano N.A., Peters G.J., Abramson R.D., Mullahy S.J.,

RA Bosak S.A., McEwan P.J., McKernan K.J., Malek J.A., Gunaratne P.H.,

RA Bosak S.A., Morley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,

RA Villalon D.K., Muzny D.M., Sodergren E.J., Lu X., Gibbs R.A.,

RA Villalon B., Ketteman M., Madan A., Godrigues S., Sanchez A.,

RA Pahey J., Helton E., Ketteman M., Madan A., Rodrigues S., Sanchez A.,

RA Hiting M., Madan A., Young A.C., Shevchenko Y., Bouffard G.G.,

RA Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M., Butterfield Y.S.,

RA Krzywinski M.I., Skalska U., Smallus D.E., Schnerch A., Schein J.E.,

RA Jones S.J., Marra M.A.,
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Best Local
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PIR; 834083; 834083.

PIR; 834086; 834086.

HSSP; P01607; 1BWW.

InterPro; IPR007110; Ig-like.

InterPro; IPR007110; Ig-v.

Pfam; PP00047; 19; 1.

SMART; SM00406; IGv; 1.

PR0SITE; P850835; IG_LIKE; 1.
                             Strausberg R.;
Submitted (JUN-2004) to the E
EMBL; BC073775; AAH73775.1; -
InterPro; IPR003599; Ig.
InterPro; IPR007110; Ig-like.
InterPro; IPR003597; Ig_c1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Q6GMX0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Hypothetical protein.
Homo sapiens (Human).
Eukaryota; Metazoa; Cl
Mammalia; Eutheria; P.
                                                                                                                                                                                                    Jones S.J., Marra M.A.; "Generation and initial analysis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              05-JUL-2004 (TrEMBLrel.
05-JUL-2004 (TrEMBLrel.
05-JUL-2004 (TrEMBLrel.
                                                                                                                                                                                        and
                                                                                                                         rissum-spleen,
                                                                                                                                           SEQUENCE
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                                                                                                                                                                       sequences.";
ad. Sci. U.S.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Chordata;
Primates;
Ig.
Ig-like.
Ig_cl.
Ig_MHC.
Ig_v.
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27,
27,
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Last sequ
Last anno
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Pred. No. 9.1e-37;
7; Mismatches 7
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                                                                                                                                                                        99:16899-16903 (2002)
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annotation update)
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                                                                                                                                                                                                                                                                                                             RX Strausberg R.L., Feingold B.A., Grouse L.H., Derge J.G.,
RA Strausberg R.L., Collins F.S., Wagner L., Shenmen C.M., Schuler G.D.,
RA Klausner R.D., Collins F.S., Wagner L., Shenmen C.M., Schuler G.D.,
RA Klausner R.D., Collins F.S., Wagner L., Shaefer C.F., Bhat N.K.,
RA Altschul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,
RA Altschenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,
RA Diatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,
RA Brownstein M.J., Ugdin T.B., Toshiyuki S., Carninci P., Prange C.,
RA Brownstein M.J., Ugdin T.B., Toshiyuki S., Carninci P., Prange C.,
RA Brownstein M.J., McKernan K.J., Malek J.A., Gunaratne P.H.,
RA B.S., Loquellano N.A., Peters G.J., Abramson R.D., Mullahy S.J.,
RA Bosak S.A., McEwan P.J., McKernan K.J., Malek J.A., Gunaratne P.H.,
RA Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,
RA Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,
RA Richards S., Worley K.C., Hale S., Garcia A.M., Gibbs R.A.,
RA Willalon D.K., Muzny D.M., Sodergren B.J., Lu X., Gibbs R.A.,
RA Wilting M., Madan A., Young A.C., Shevchenko Y., Bouffard G.G.,
RA Whiting M., Madan A., Young A.C., Shevchenko Y., Bouffard G.G.,
RA Wilting M., Madan A., Young A.C., Shevchenko Y., Boutfard G.G.,
RA Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M., Butterfield Y.S.,
RA Krzywinski M.I., Skalska U., Smailus D.B., Schnerch A., Schein J.B.,
Ra Krzywinski M.I., Skalska U., Smailus D.B., Schnerch A., Schein J.B.,
Ra Krzywinski M.I., Skalska U., Smailus D.B., Schnerch A., Schein J.B.,
Ra Krzywinski M. I., Skalska U., Smailus D.B., Schnerch A., Schein J.B.,
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Pfam; PP07654; C1-set; 1.
Pfam; PP00047; ig; 2.
SMART; SM00409; IG; 2.
SMART; SM00407; IGc1; 1.
SMART; SM00406; IGV; 1.
SMART; SM00406; IGV; 1.
PROSITE; P850835; IG_LIKE; 2.
PROSITE; P8508290; IG_MHC; UNKNOWN_1.
                Strausberg R.;
Strausberg R.;
Submitted (JUN-2004) to the EMBL/GenBank/DDBJ databases.
EMBL, BC073791; AAH73791.1; -.
InterPro; IPR003599; Ig.
InterPro; IPR003199; Ig.-like.
InterPro; IPR003597; Ig.-c1.
InterPro; IPR003087; Ig.MHC.
InterPro; IPR00306; Ig.MHC.
InterPro; IPR00306; Ig.MHC.
InterPro; IPR00306; Ig.MHC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Q6GMW1;
Q6GMW1;
05-JUL-2004
05-JUL-2004
05-JUL-2004
                                                                                                                                                                                             SEQUENCE FROM
TISSUE=Spleen;
                                                                                                                                                                                                                                                                   and
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Mammalia; Eutheria; Primates;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               GVPSRFSGSGSGTDFTLTISSLRPDDFATYYCQQSYNIPLTFGGGTNVEIKRTV
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l. Acad. Sci. U.S.
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19; 2.
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Pred. No. 3.1e-36;
2; Mismatches 9;
                                                                                                                                                                                                                                                  99:16899-16903 (2002)
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Catarrhini; Hominidae;
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annotation update)
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WEDLINE-22388257; PubMed=12477932;

A Strausberg R.L., Feingold E.A., Grouse L.H., Derge J.G.,

Klausner R.D., Collins F.S., Wagner L., Shenmen C.M., Schuler G.D.,

A Altschall S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,

A Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,

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A Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,

A Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,

A Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,

A Hopkins R.F., Jordan R.B., Bonaldo M.F., Casavant T.L., Scheetz T.E.,

Brownstein M.J., Usdin T.B., Toshiyuki S., Carninci P., Frange C.,

A Brownstein M.J., Usdin T.B., Toshiyuki S., Carninci P., Frange C.,

A Brownstein M.J., Wedlin T.B., Toshiyuki S., Carninci P., Frange C.,

A Bosak S.A., McZwan P.J., McKernan K.J., Malek J.A., Gunaratne P.H.,

A Richards S., Morley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,

Villalon D.K., Muzny D.M., Sodergren B.J., Lu X., Gibbs R.A.,

A Villalon D.K., Muzny D.M., Sodergren B.J., Lu X., Gibbs R.A.,

A Fahey J., Helton B., Ketteman M., Madan A., Rodrigues S., Sanchez A.,

Whiting M., Madan A., Young A.C., Shevchenko Y., Bouffard G.G.,

Blakesley R.W., Touchman J.W., Green B.D., Dickson M.C.,

A Krzywinski M.I., Skalska U., Smailus D.E., Schnerch A., Schein J.E.,

Jones S. T. Mayrra M.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
Best Local
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                         InterPro; IPR003599; Ig.
InterPro; IPR007110; Ig-like.
InterPro; IPR00710; Ig-like.
InterPro; IPR003597; Ig_cl.
InterPro; IPR003006; Ig_MHC.
InterPro; IPR003596; Ig_v.
Pfam; PP07654; Cl-set; I.
Pfam; PF070654; Cl-set; I.
SMART; SM00407; Ig; 2.
SMART; SM00406; IGv; 1.
SMART; SM00406; IGv; 1.
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SMART; SM00406; IGv; 1.
PROSITE; PS50835; IG_LIKE; 2.
PROSITE; PS00290; IG_MHC; UNKNOWN
Hypothetical protein.
SEQUENCE 236 AA; 25751 MW; 5BI
                                                                                                                                                                                                                                                                                                                                     Strausberg R.;
Submitted (JUL-2002) to the
Submit BC034141; AAH34141.1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Jones S.J., Marra M.A.;
"Generation and initial analysis of more than 15,000 full-length and mouse CDNA sequences.";
and mouse CDNA sequences.";
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Homo sapiens (Human).
Bukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Primates;
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                                                                                                                                                                                                                                                                                                                                                                                                                                  TISSUE=Lung;
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05-JUL-2004
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sci.
   IG_LIKE;
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Last sequence update)
Last annotation updat
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Pred. No. 3.9e-36;
6; Mismatches 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Craniata; Vertebrata; Euteleostomi; Catarrhini; Hominidae; Homo.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  236
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RA Klausner R.D., Collins F.S., Wagner L., Shenmen C.M., Schuler G.D.,
RA Altschul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,
RA Altschul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,
RA Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Heleh F.,
RA Stapleton M., Soares M.B., Bonaldo M.F., Casavant T.L., Scheetz T.E.,
Stapleton M., Soares M.B., Bonaldo M.F., Casavant T.L., Scheetz T.E.,
RA Stapleton M., Soares M.B., Bonaldo M.F., Casavant T.L., Scheetz T.E.,
RA Stapleton M., Soares M.B., Bonaldo M.F., Casavant T.L., Scheetz T.E.,
RA Stapleton M., Soares M.B., Bonaldo M.F., Casavant T.L., Scheetz T.E.,
RA Stapleton M., Soares M.B., Bonaldo M.F., Casavant T.L., Scheetz T.E.,
RA Brownstein M.J., Ugdin T.B., Toshiyuki S., Carninci P., Prange C.,
RA Raha S.S., Loquellano N.A., Peters G.J., Abramson R.D., Mullahy S.J.,
RA Raha S.S., Loquellano N.A., Peters G.J., Abramson R.D., Mullahy S.J.,
RA Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,
RA Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,
RA Villalon D.K., Muzny D.M., Sodergren B.J., Lu X., Gibbs R.A.,
RA Wilting M., Madan A., Young A.C., Shevchenko Y., Bouffard G.G.,
RA Whiting M., Madan A., Young A.C., Shevchenko Y., Bouffard G.G.,
RA Modriguez A.C., Grimwood J., Schmutz J., Myers R.M., Butterfield Y.S.,
RA Krzywinski M.I., Skalska U., Smailus D.B., Schnerch A., Schein J.B.,
RA Krzywinski M.I., Skalska U., Smailus D.B., Schnerch A., Schein J.B.,
                                                                                                                              Query Match
Best Local S
Matches 90
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Best Local (
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02-MAR-2004
02-MAR-2004
02-MAR-2004
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Hypothetical protein.

SEQUENCE 236 AA; 25603 MW; BBC56
                                                                                                                                                                                                                              Strausberg R.;
submitted (JUL-2002) to the
EMBL; BC034141; AAH34141.1;
                                                                                                                                                                                                  Hypothetical protein. SEQUENCE 236 AA; 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TISSUE=Lung;
MEDLINE=22388257; PubMed=12477932;
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Mammalia; Eutheria;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Hypothetical protein. Homo sapiens (Human).
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                                                                                                                                                                                                                                                                                rissum=Lung;
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                                                                                                                                                                                                                                                                                                   EQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                 S.J., Marra M.A.;
                             61
                                                              23
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79
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                                                         DIQMTQSPSSLSASVGDRVTITCRASQSVSTSSYSYMHWYQQKPGKAPKVLISYASNLES
             GVPSRFSGSGSGTDFTLTISSLQPEDFATYYCQHSWGIPRTFGQGTKVEIKRTV
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(TrEMBLrel.
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Primates;
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78.9%;
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78.98;
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Pred. No. 3.7e-35;
                                                                                                                                                Score 444; DB 2;
Pred. No. 3.7e-35;
                                                                                                                                                                                                                                                                                                                                    99:16899-16903 (2002).
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annotation update)
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RESULT 10
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AC Q9UL7
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DT 01-OC
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RN [1] -
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RA WG X.
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Best Local &
Matches 87
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HSSP; P01607; 1BWW.
GO: GO:0005576; C:extracellular; NAS.
GO: GO:0003823; F:antigen binding; NAS.
GO: GO:0006955; P:immune response; NAS.
InterPro; IPR007110; Ig-like.
InterPro; IPR003596; Ig_v.
[1]
SEQUENCE FROM N.A.
MEDLINE=98277139; PubMed=9614934;
rin B., Van der Merwe P.L.
                                                                                                                                             Q9UL70;
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DISULFID
NON TER
SEQUENCE
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01-MAY-2000 (TrEMBLrel.
01-OCT-2003 (TrEMBLrel.
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SMART; SM00406; IGv; 1.
PROSITE; PS50835; IG_LIKE;
Direct protein sequencing;
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                                             Eukaryota; Metazoa;
Mammalia; Eutheria;
NCBI_TaxID=9606;
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21-JUL-1986 (Rel.
05-JUL-2004 (Rel.
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                                                                                                   Myosin-reactive
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Homo
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                                                                               (Fragment).
Homo sapiens (Human)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           - MISCELLANEOUS:
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amino acid sequence of a human kappa ligh
hem. J. 123:945-958(1971).
MISCELLANEOUS: The C region of this chain
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               HUMAN
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EMBLrel. 13, Last sequence update)
EMBLrel. 25, Last annotation update)
immunoglobulin light chain variable
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                                                        Chordata;
Primates;
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49
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77.7%;
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Pred. No. 3e-35;
9; Mismatches ]
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                                                           Craniata; Vo
Catarrhini;
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         Kalis
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                                                                    Vertebrata;
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        Berney S.M.,
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                                                                       Euteleostomi;
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Best Local S
Matches 89
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P01606;
21-JUL-1986
21-JUL-1986
05-JUL-2004
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CMART: SM00406; IGV; 1.
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InterPro; IPR003596; Ig_v.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Eukaryota; Metazoa;
Mammalia; Eutheria;
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                                                                                                                                                                                                   PROSITE;
                                                                                                                                                                                                                                                                                                                                                                                                       -!- MISCELLANEOUS: The C regic
                                                                                                                                                                                                                                                                                                                                                                                                                                            Science 169:56-59(1970)
                                                                                                                                                                                                                                                                                                                                                                                                                                                               chains."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NCBI_TaxID=9606;
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SMART; SM00406; IGv;
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GO:0005576; C:extracellular; NAS.
GO:0003823; F:antigen binding; NA
GO:0006955; P:immune response; NA
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                                                                                                                                                                               protein
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                                  n sequencing;
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24 34
35 49
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98 107
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Primates;
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G_LIKE;
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79.5%;
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X
X
                              Framework-2.
Complementarity-determining-2.
Framework-3.
Complementarity-determining-3.
Framework-4.
By similarity.
                                                                                                                                                                               Immunoglobulin
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Pred. No. 3e-3
5; Mismatches
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                                                                                                                                           Complementarity-determining-1.
                                                                                                                                                              Framework-1
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Catarrhini; Hominidae;
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8283D4A24105827E
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RESULT 12
KV1W_HUMAN
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Best Local S
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DOMAIN
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SMART; SM00406; IGv; 1.
PROSITE; PS50835; IG LIKE;
                                                                                                                                                                                                                                                                                                                                                     EMBL; X00965; CAA25477.1; PIR; A01883; K1HUWK.
                                                                                                                                                                                                                                                                                                                                                                                       or send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                                                                   entities
                                                                                                                                                                                                                                                                                                                                                                                                            use by non-profit institutions as long modified and this statement is not removed.
                                                                                                                                                                                                                                                                                                                                                                                                                          the European Bioinformatics Institute. There are no resume by non-profit institutions as long as its content
                                                                                                                                                                                                                                                                                                                                                                                                                                  This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation the European Bioinformatics Institute. There are no restrictions on its
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Klobeck H.G., Combriato G., Zachau H.G.; Immunoglobulin genes of the kappa light cl lymphoid cell lines are closely related."; Nucleic Acids Res. 12:6995-7006(1984).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SEQUENCE FROM N.A. MEDLINE=85014148; PubMed=6091049;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; I
Mammalia; Eutheria; Primates; Catarrhini; Hominidae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Homo sapiens (Human)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              P04431;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          KV1W
                                                                                            SEQUENCE
                                                                                                                                                                                                                                                                            InterPro; IPR007110; Ig-like.
InterPro; IPR003596; Ig_v.
                                                                                                                                                                                                                                                                                                                                           HSSP;
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                                                                                                                              DOMAIN
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-AUG-1987 (Rel. 05, Last sequence update)
-UUL-1999 (Rel. 38, Last annotation update)
kappa chain V-I region Walker precursor.
                                                                                                                                                                                                                                                                                                    GO:0005576; C:extracellular; NAS.
GO:0003823; F:antigen binding; NF
GO:0006955; P:immune response; NF
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                                                                                                                                                                                                                                                                                                                                           P01607; 1BWW.
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               DIQMTQSPSSLSASVGDRVTITCRASQSVSTSSYSYMHWYQQKPGKAPKVLISYASNLES
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                                               Conservative
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69.68;
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79.3%;
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                                                                                            MW;
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                                              10;
                                                                                                                           Ig kappa chain V-I region Walker
Framework-1.
Complementarity-determining-1.
Framework-2.
Complementarity-determining-2.
Framework-3.
Complementarity-determining-3.
Framework-4.
                                           Score 440; DB 1; 1
Pred. No. 4.6e-35;
0; Mismatches 9;
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Pred. No. 3.7e-35;
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                                                                                                                                                                                                           kappa chain V-I region Walker.
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RA Allausner R.D., Collins F.S., Wagner L., Schaefer C.F., Bhat N.K.,
RA Altachul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,
RA Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,
RA Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,
RA Stapleton M., Soares M.B., Bonaldo M.F., Casavant T.L., Scheetz T.E.,
RA Stapleton M.J., Usdin T.B., Toshiyuki S., Carninci P., Prange C.,
RA Brownstein M.J., Usdin T.B., Toshiyuki S., Carninci P., Prange C.,
RA Brownstein M.J., Wachin T.B., Toshiyuki S., Carninci P., Prange C.,
RA Brownstein M.J., Wachin T.B., Toshiyuki S., Carninci P., Prange C.,
RA Raha S.S., Loquellano N.A., Peters G.J., Abramson R.D., Mullahy S.J.,
RA Robak S.A., McEwan P.J., McKernan K.J., Malek J.A., Gunaratne P.H.,
RA Bosak S.A., McEwan P.J., McKernan K.J., Lu X., Gibbs R.A.,
RA Villalon D.K., Muzny D.M., Sodergren E.J., Lu X., Gibbs R.A.,
RA Villalon D.K., Muzny D.M., Sodergren E.J., Lu X., Gibbs R.A.,
RA Whiting M., Madan A., Young A.C., Shevchenko Y., Bouffard G.G.,
RA Blakesley R.W., Touchman J.W., Green E.D., Dickson M.C.,
RA Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M., Butterfield Y.S.,
RA A Tones S. T. Marra M.A.,
RA A. Tones S. T. Marra M.A.,
                                                                                                                                   Matches
                                                                                                                                                                    Query Match
                                                                                                                                                                                                                                                                   InterPro; IPR007110; Ig-like.
InterPro; IPR003597; Ig_cl.
InterPro; IPR003597; Ig_wHC.
InterPro; IPR003596; Ig_w.
Pfam; PF07654; Cl-set; 1.
Pfam; PF00047; Ig; 1.
SMART; SM00406; IGv; 1.
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01-OCT-2003
01-OCT-2003
                                                                                                                                                                                                                                                                                      Pfam;
Pfam;
                                                                                                                                                                                                                                                                                                                                                                                    TISSUE-Skeletal Muscie;
Strausberg R.;
Strausberg R.;
Submitted (MAR-2001) to the EMBL/GenBank/DDBJ databases
                                                                                                                                                                                                     Hypothetical SEQUENCE 2:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SEQUENCE FROM N.A.
TISSUE=Skeletal Muscle;
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Mammalia; Eutheria;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Hypothetical protein.
Homo sapiens (Human).
                                                                                                                                                                                                                                  PROSITE; PS50835; IG_LIKE; 2.
PROSITE; PS00290; IG_MHC; UNKNOWN_1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        and mouse cDNA sequences.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NCBI_TaxID=9606;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Hypothetical
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       "Generation and initial analysis of more than 15,000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Jones S.J., Marra M.A.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          MEDLINE=22388257; PubMed=12477932;
                                                                                                                                                                                                                                                                                                                                                                                                                                                       EQUENCE FROM N.A.
                                                                                                                                                    Local
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 79
                                                                                                  μ
                                                                                                                                   99;
                                                                                                                                                    Similarity
                      GVPSRFSGSGSGTDFTLTISSLQPEDFATYYCQHSWGIPRTFGQGTKVEIKRTV 114
                                                                                        DIQMTQSPSSLSASVGDRVTITCRASQSVSTSSYSYMHWYQQKPGKAPKVLISYASNLES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   GVTSRFSGSGSGTDFTLTISSLQPEDSATYYCQQSYSTLITFGQGTRLEIK 129
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GVQSKFSGSGSGTDFTLTISSLQPEDFATYYCQQYKSYPVTFGQGTKLEIKRTV
                                                                   DIQMTQSPSSLSASVGDTVTITCRASQDIS----NYLAWFQQKPGKAPKSLIYGASSLQS
                                                                                                                                                                                                     al protein.
236 AA; 2
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(TrEMBLrel.
(TrEMBLrel.
                                                                                                                                   Conservative
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Primates;
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78.1%;
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Last sequence update)
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                                                                                                                                   8
                                                                                                                                                    Score 440; DB 2;
Pred. No. 9.1e-35;
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Catarrhini;
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                                                                                                                                                                                                     7FBFE4ED23084BC6 CRC64;
                                                                                                                                     Mismatches
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RESULT 14

RYIK TUTYAN

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AC PO160

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PO 21-JU

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Best Local
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21-JUL-1986 (Rel. 01,
21-JUL-1986 (Rel. 01,
01-OCT-2004 (Rel. 45
01-OT-2004 (Rel. 45
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   KV1K HUMAN
P01603;
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                                                                                                                                                                                                                                                                          RTAX
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NON TER
SEQUENCE.
MEDLINE=72189444; PubMed=5028201;
Schiechl H., Hilschmann N.;
                                                                                 Eukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Primates;
MCBI_TaxID=9606;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             DOMAIN
DOMAIN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DOMAIN
DOMAIN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   GO; GO:0005576; C:extracellular; NAS. GO; GO:0003823; F:antigen binding; NAS. GO; GO:0008955; P:immune response; NAS. InterPro; IPR007110; Ig-11ke. InterPro; IPR003596; Ig_v.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  "Comparative structural studies on the light chains of human immunoglobuling. I. Protein Ka with the Inv(3) allotypic marker.";
J. Blochem. 77:1277-1296(1975).
-I- MISCELLANEOUS: The C region of this chain has the INV (3) marker.
-I- MISCELLANEOUS: This is a Bence-Jones protein.
                                                                                                                                            Ig kappa chain V-I r
Homo sapiens (Human)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Pfam; PF00047; 1g; 1.
SMART; SM00406; IGv;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PIR; A01869; K1HUKA.
HSSP; P80362; 1WTL.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            DOMAIN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Bence-Jones
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PROSITE; PS50835;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Shinoda
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           MEDLINE=76189985;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NCBI_TaxID=9606,
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350835; IG_LIKE; 1.
3 protein; Direct pr
1lin V region.
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44, Last
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01, Last sequence update)
45, Last annotation update)
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73.2%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      protein sequencing;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Score 438;
Pred. No. 5.
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By similarity.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Complementarity-determining-2
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                                                                                                          Craniata; Vertebrata; Euteleostomi; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Framework-1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PRT;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              768839FBED5A2F4B CRC64;
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on update)
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Best Local S
Matches 88
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DOMAIN 24
DOMAIN 35
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DOMAIN 50
DOMAIN 57
DOMAIN 89
DOMAIN 98
DISULFID 23
STRAND 10
STRAND 10
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                                                                                                                                                                                                                                                                                                             TURN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               9998
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PROSITE; PS50835; IG LIKE; 3D-structure; Bence-Jones;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SMART; SM00406; IGv; 1.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            molecular replacement methods
region of the kappa chain RBI.
-I- MISCELLANEOUS: The C region of
-I- MISCELLANEOUS: This is a Bence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Fehlhammer H., Schiffer M., Epp O., Schwager P., Steigemann W., Schramm
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              X-RAY CRYSTALLOGRAPHY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Hoppe-Seyler's Z.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Jones protein Au."
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           InterPro; IPR007110; Ig-like.
InterPro; IPR003596; Ig_v.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The structure determination of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ; 1JV5; X-ray; A=1-107.
GO:0005576; C:extracellular; NAS.
GO:0003823; F:antigen binding; NAS.
GO:0006955; P:immune response; NAS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           phys. Struct. Mech. 1:139-146(1975).
MISCELLANEOUS: The structure of the V region of the V reg
 57
                             61
                                                                                                                                88;
                                                                                               \vdash
                                                                                                                                               Similarity
                             GVPSRFSGSGSGTDFTLTISSLQPEDFATYYCQHSWGIPRTFGQGTKVBIKR
                                                             DIQMTQSPSSLSASVGDRVTITCQASQDIS----DYLNWYQQKPGKAPKLLIYDASNLES
                                                                                              DIOMTOSPSSLSASVGDRVTITCRASQSVSTSSYSYMHWYQQKPGKAPKVLISYASNLES
GVPSRFSGGGSGAHFTFT
                                                                                                                                Conservative
                                                                                                                                                                                                                                                                                               AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           V region.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Physiol. Chem. 353:345-370(1972)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PubMed=1234024;
                                                                                                                                                                                                                              11939 MW;
                                                                                                                                             73.6%;
78.6%;
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                                                                                                                              Score 437; DB 1;
Pred. No. 7.3e-35;
6; Mismatches 14
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                                                                                                                                                                                               E8011187EE6F6FB9 CRC64;
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IATYYCQQYDYLPWTFGQGTKVEIKR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Colman P.M., Lattman H.J.; variable portion of 1
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                                                                                                                              14;
                                                                                                                                                             Length 108;
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Job time : 128.21 secs

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Result
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Maximum Match 100%
Listing first 45 summaries
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Minimum DB seq length: 0
Maximum DB seq length: 2000000000
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221098765422
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                                                                                                                                                                        Score
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                                                                                                                                                                                     Query
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                                                                                                                                                                                                                                                                                      Published_Applications_AA: *
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Copyright (c) 1993 - 2004 Compugen Ltd.
                                                                                                                                                                         Length
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 1 14 US-10-044-896-3

15 US-10-378-567-3

9 US-09-917-410-2

14 US-10-216-484-127

14 US-10-384-933-127

14 US-10-384-933-127

14 US-10-384-933-131

14 US-10-384-933-131

19 US-09-920-171-10

10 US-10-113-996-10

10 US-10-113-996-10

10 US-10-791-619-10

9 US-09-802-077-9

9 US-09-802-096-9

9 US-09-920-171-13
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Sequence 3, Appli
Sequence 2, Appli
Sequence 27, App
Sequence 127, App
Sequence 131, App
Sequence 131, App
Sequence 131, App
Sequence 10, Appl
Sequence 10, Appl
Sequence 10, Appl
Sequence 9, Appli
Sequence 9, Appli
Sequence 9, Appli
Sequence 9, Appli
Sequence 13, Appli
Sequence 13, Appli
                                                                                                                                                                      Description
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Sequence 15		10	110	80.4	
	US-09-974-051-	φ	110	80.4	477.5
	US-09-974-052-1	9	110	•	•
	US-09-025-403A-15	ø	110	80.4	٠
	US-10-044-896-	14	114	80.5	478
	US-09-974-516-	10	110	•	480.5
	US-09-974-498	10	110	80.9	480.5
		9	110	•	480.5
	US-09-974-052-1	9	110	•	480.5
	ď	9	110	•	480.5
	-Sn	17	111	81.0	481
	Sn	14	111	•	481
	ď	9	111	81.0	481
		17	108		482
		17	240	81.3	483
	us-	11	240	•	483
	US-09-968-744A	10	240	•	483
	US-09-968-561A	9	240	81.3	483
	ď	ø	240	•	483
	US-10-791-619-	17	114	٠	483
		14	114	•	483
	ď	9	114	٠	483
	US-10-409-814A	17	108	81.3	483
	US-10-663-244	16	233	٠	484
	-10-384-933-12	14	238	•	485
	US-10-216	14	238	81.6	485
	US-10-764-42	17	669		486
	US-10-764-428-21	17	669	•	486
	US-10-791-61	17	218		486
	-10-813-483-	17	218		486
	US-10-113-996-13	14	218	81.8	486
	US-09-925-179-9	10	218	81.8	486

### ALIGNMENTS

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GENERAL INFORMATION:

APPLICANT: Chuntharapai, Anan

APPLICANT: Kim, Jin K.

APPLICANT: Kiewart, Timothy

APPLICANT: Presta, Leonard G.

TITLE OF INVENTION: ANTI-INVERFERON-ALPHA ANTIBODIES

FILE REFERENCE: GENENT: 0.74A

CURRENT APPLICATION NUMBER: US/10/044,896

CURRENT FILING DATE: 2002-01-09

PRIOR APPLICATION NUMBER: 60/270775

PRIOR FILING DATE: 2001-02-22

NUMBER OF SEQ ID NOS: 14

SOFTWARE: FastSEQ for Windows Version 4.0

SEQ ID NO 3

LENGTH: 114

TYPE: PRT
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                                                                                                                                                                                                                                                                  US-10-044-896-3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       RESULT 1
US-10-044-896-3
                                                                                                                                                                  Query Match 100.0%; Score 594; DB 14; Best Local Similarity 100.0%; Pred. No. 1.8e-40; Matches 114; Conservative 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 3, Application US/10044896 Publication No. US20030166228A1
                                                                                                                                                                                                                                                                               ORGANISM: Artificial Sequence FEATURE:
OTHER INFORMATION: This sequence represents a humanized chimeric OTHER INFORMATION: antibody comprising human and non-human sequences.
61 GVPSRFSGSGGTDFTLTISSLQPEDFATYYCQHSWGIPRTFGQGTKVEIKRTV 114
                                                                     DIQMTQSPSSLSASVGDRVTITCRASQSVSTSSYSYMHWYQQKPGKAPKVLISYASNLES
                                                                                                                                                                                                               Length 114;
                                                                                                                                                                Indels 0;
                                                                                                                                                                  Gaps
                                                                                                                        60
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GVPSRFSGSGSGTDFTLTISSLQPEDFATYYCQHSWGIPRTFGQGTKVBIKRTV 114

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RESULT 3
US-09-917-410-2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                US-10-378-567-3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 2, Application US/09917410 Patent No. US20020098183A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SEQ ID NO 3
LENGTH: 218
TYPE: PRT
ORGANISM: Artificial Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PRIOR APPLICATION NUMBER: 60/276,452
PRIOR FILING DATE: 2001-03-16
PRIOR APPLICATION NUMBER: 60/229,933
PRIOR FILING DATE: 2000-09-01
NUMBER OF SEQ ID NOS: 3
SOFTWARE: Patentin Ver. 2.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CURRENT APPLICATION NUMBER: US/10/378,567
CURRENT FILING DATE: 2003-02-28
PRIOR APPLICATION NUMBER: PCT/US01/27352
PRIOR FILING DATE: 2001-08-31
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TITLE OF INVENTION: CO-CRYSTAL STRUCTURE OF MONOCLONAL ANTIBODY 5C8 AND TITLE OF INVENTION: CD154, AND USE THEREOF IN DRUG DESIGN FILE REFERENCE: A096CON1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: KARPUSAS, APPLICANT: HSU, YEN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                OTHER INFORMATION: Description of Artificial Sequence: Synthetic OTHER INFORMATION: humanized 5c8 light chain amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Local Similarity
                     COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS SOFTWARE: ASCII, WordPerfect 5.1 CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                    TITLE OF INVENTION: ANTI-L-SELECTIN ANTIBODIES FOR PREVENTION OF MULTIPLE ORGAN FAILURE AFTER POLYTRAUMA AND FOR PREVENTION OF ACUTE ORGAN DAMAGE AFTER
                                                                                                           COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Computer Disk
                                                                                                                                                                                                                                                                              CORRESPONDENCE ADDRESS:
ADDRESSEE: Felfe & Lynch
                                                                                                                                                                                                                                                                                                                           NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: MARTIN, Ulrich; HASELBECK, Anton; SCHUMACHER, Guenther;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    61
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          61 GVPSRFSGSGSGTDFTLTISSLQPEDFATYYCQHSWGIPRTFGQGTKVEIKRTV 114
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   91;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1 DIVLTQSPATLSVSPGERATISCRASQRVSSSTYSYMHWYQQKPGQPPKLLIKYASNLES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1 DIOMTOSPSSLSASVGDRVTITCRASOSVSTSSYSYMHWYQOKPGKAPKVLISYASNLES
                                                                                                                                                                                                        STREET: 805 Third Avenue
CITY: New York
STATE: New York
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           GVPARFSGSGSTDFTLTISSVEPEDFATYYCQHSWEIPFTFGGGTKLEIKRTV 114
APPLICATION NUMBER: US/09/917,410
                                                                                                                                                                                   COUNTRY: U.S.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TAYLOR, FREDERICK R. ZHENG, ZHONGLI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Conservative
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YEN-MING
                                                                                                                                                                                                                                                                                                                                                                                                                                           CO, Man S.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   83.2%; Score 494; DB 15; 79.8%; Pred. No. 3.7e-32; tive 13; Mismatches 10;
                                                                                                                                                                                                                                                                                                                                                EXTRACORPOREAL BLOOD CIRCULATION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     10;
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Sequence 127, Application US/10216484

Publication No. US20030103976A1

GENERAL INFORMATION:
APPLICANT: Haruyama, No. US20030103976A1ufusa
APPLICANT: Haruyama, Hideyuki
APPLICANT: Haruyama, Kaori
APPLICANT: Tamaki, Ituko
APPLICANT: Tamaki, Ituko
APPLICANT: Tamaki, Tohru
ITILE OF INVENTION: Anti-Fas Antibodies
FILS REFERENCE: 980126CIP/HG
CURRENT APPLICATION NUMBER: US/10/216,484
CURRENT APPLICATION NUMBER: US/09/499,662
PRIOR APPLICATION NUMBER: US/09/499,662
PRIOR FILING DATE: 2000-02-09
PRIOR FILING DATE: 1908-04-01

PRIOR APPLICATION NUMBER: US 09/053,583
PRIOR APPLICATION NUMBER: US 09/053,583
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                                                                                                                                                                 ; OTHER INFORMATION: Description of Artificial Sequence: Designed light; OTHER INFORMATION: chain of humanized anti-Fas antibody US-10-216-484-127
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                                                                                   Matches
                                                                                                                                                                                                                                                                                                                    NUMBER OF SEQ ID NOS: 165
SEQ ID NO 127
                                                                                                         Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Matches
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TELEPHONE: (212) 688-9200
TELEPAX: (212) 838-3884
INFORMATION FOR SEQ ID NO: 2:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                     ORGANISM: Artificial Sequence FEATURE:
                                                                                                                                                                                                                                                                             TYPE: PRT
                                                                                                                                                                                                                                                                                                   LENGTH: 238
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#ILING DATE: <UNKNOWN>
APPLICATION UNMBER: EP 95 114 969.9

#ILING DATE: 19-6ep-95

#ITING DATE: 19-6ep-95

#ITORNEY/AGENT IMFORMATION:
NAME: Hanson, No. US20020098183Alman D.
### REGISTIATION NUMBER: 30,946

### REFERENCE/DOCKET NUMBER: BOER 1059-PFF/NDH/SLH
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TOPOLOGY: linear MOLECULE TYPE: protein SEQUENCE DESCRIPTION: SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              FILING DATE: 26-Jul-2001
CLASSIFICATION: <Unknown
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              98;
  21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     61 GVPSRFSGSGSGTDFTLTISSLQPEDFATYYCQHSWGIPRTFGQGTKVEIKRTV 114
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             61 GIPSRFSGSGSGTDFTLTISSLQPEDFATYYCQQSNEDPWTFGQGTKVEIKRTV 114
                              1 DIQMTQSPSSLSASVGDRVTITCRASQSVSTSSYSYMHWYQQKPGKAPKVLISYASNLES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1 DIQMTQSPSSLSASVGDRVTITCRASQSVSTSSYSYMHWYQQKPGKAPKVLISYASNLES
                                                                                                         Similarity
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DIVLTQSPSSLSASVGDRVTITCKASQSVDYDGDSYMNWYQQKPGKAPKLLIYAASNLES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               DIQMTQSPSSLSASVGDRVTITCKASQSVDYDGDSYMNWYQQKPGKAPKLLIYAASNLES
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                                                                                                   Score 489; DB 1
Pred. No. 1e-31;
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Pred. No. 7.8e-32;
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                                                                                                                         DB 14;
                                                                                      12;
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APPLICANT: Tamaki, ikuko

APPLICANT: Takahashi, Tohru

FIILE OP INVENTION: Anti-Fas Antibodies

FILE REFERENCE: 980126CIP/HG

CURRENT APPLICATION NUMBER: US/10/216,484

CURRENT FILING DATE: 2000-08-09

PRIOR APPLICATION NUMBER: US/09/499,662

PRIOR FILING DATE: 2000-02-09

PRIOR APPLICATION NUMBER: US 09/053,583

PRIOR FILING DATE: 1998-04-01

NUMBER OF SEQ ID NOS: 165

SEQ ID NO 131

LENGTH: 77
, OTHER INFORMATION: Description of Artificial Sequence: Designed light, OTHER INFORMATION: chain of humanized anti-Fas antibody US-10-216-484-131
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    US-10-216-484-131
; Sequence 131, Application US/10216484
; Publication No. US20030103976A1
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US-10-384-933-127
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SEQ ID NO 127
LENGTH: 238
TYPE: PRT
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CURRENT FILING DATE: 2003-02-05
PRIOR APPLICATION NUMBER: US/09/499,662
PRIOR FILING DATE: 2000-02-09
PRIOR APPLICATION NUMBER: EARLIER APPLICATION NUMBER: US 09/053,583
PRIOR FILING DATE: EARLIER FILING DATE: 1998-04-01
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: Haruyama, Hideyuki
APPLICANT: Nakahara, Kaori
APPLICANT: Tamaki, Ikuko
APPLICANT: Takahashi, Tohru
TITLE OF INVENTION: Anti-Fas Antibodies
FILE REFERENCE: 980126CIP/HG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Serizawa, No. US20030170817Alufusa
APPLICANT: Haruyama, Hideyuki
APPLICANT: Nakahara, Kaori
                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Serizawa, No. US20030103976Alufusa
APPLICANT: Haruyama, Hideyuki
APPLICANT: Nakahara, Kaori
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     OTHER INFORMATION: Description of Artificial Sequence: Designed light OTHER INFORMATION: chain of humanized anti-Fas antibody
                                                                          ORGANISM: Artificial Sequence FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     FEATURE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         81
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     61 GVPSRFSGSGSGTDFTLTISSLQPEDFATYYCQHSWGIPRTFGQGTKVEIKRTV 114
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Pred. No. 1e-31;
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                                                                                                                                                                                                                                                                                                                                                                      RESULT 8
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                                                                                                                                                                                              GENERAL INFORMATION:
APPLICANT: Lowman, Henry B.
APPLICANT: Presta, Leonard
APPLICANT: Jardieu, Paula
APPLICANT: Lowe, John
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SEQ ID NO 131
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                                                                                                                                                                                                                                                                                                       Sequence 10, Application US/09920171 Patent No. US20020054878A1
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TITLE OF INVENTION: Improved Anti-IgE Antibodies FILE REFERENCE: F1123C2US
CURRENT EPELICATION NUMBER: US/09/920,171
CURRENT FILING DATE: 2001-08-01
PRIOR APPLICATION NUMBER: US 08/887,352
PRIOR FILING DATE: 1997-07-02
PRIOR FILING DATE: US 09/296,005
PRIOR APPLICATION NUMBER: US 09/296,005
PRIOR PILING DATE: 1999-04-21
NUMBER OF SEQ ID NOS: 44
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PRIOR FILING DATE: 2000-02-09
PRIOR APPLICATION NUMBER: US 09/053,583
PRIOR FILING DATE: EARLIER APPLICATION NUMBER: US 09/053,583
PRIOR FILING DATE: EARLIER FILING DATE: 1998-04-01
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CURRENT APPLICATION NUMBER: US/10/384,933
CURRENT FILING DATE: 2003-02-05
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TITLE OF INVENTION: Anti-Fas Antibodies
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APPLICANT: Haruyama, Hideyuki
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TYPE: PAT
TYPE: ORGANISM: Artificial Sequence
PEATURE:
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Jardieu, Paula M.
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Tamaki, Ikuko
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o. US20030170817A1
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Pred. No. 1.2e-31;
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Pred. No. 1.2e-31;
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TITUE OF INVENTION: Improved Anti-IgE Antibodies
FILE REFERENCE: p1123GUS
CURRENT APPLICATION NUMBER: US/10/113,996
CURRENT FILING DATE: 2002-04-01
PRIOR APPLICATION NUMBER: US 08/887,352
PRIOR FILING DATE: 1997-07-02
PRIOR APPLICATION NUMBER: US 09/296,005
PRIOR APPLICATION NUMBER: US 09/296,005
PRIOR APPLICATION NUMBER: US 09/296,171
PRIOR FILING DATE: 1999-04-21
PRIOR FILING DATE: 2001-08-01
PRIOR FILING DATE: 2001-08-01
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; OTHER INFORMATION: Light chain sequence derived from MAB11
US-09-920-171-10
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               Sequence 10, Application US/10791619
Publication No. US20040259077A1
GENERAL INFORMATION:
APPLICANT: Henry B. Lowman, Leonard G. Presta, Paula M. Jardieu, John Lowe
TITLE OF INVENTION: Improved Anti-IgE Antibodies and Method of Improving Polypeptides
FILE REFERENCE: P11281
CURRENT APPLICATION NUMBER: US/10/791,619
CURRENT FILING DATE: 2004-03-02
CURRENT FILING DATE: 2004-03-02
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SEQ ID NO 10
LENGTH: 114
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LENGTH: 114
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PRIOR APPLICATION NUMBER: US/09/109,207
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Pred. No. 8.8e-32
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Pred. No. 8.8e-32;
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RESULT 12
US-09-802-096-9
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PRIOR APPLICATION NUMBER: US 60,
PRIOR FILING DATE: 1997-07-03
NUMBER OF SEQ ID NOS: 44
SEQ ID NO 10
LENGTH: 114
TYPE: PRT
ORGANISM: Artificial
FEATURE:
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US-09-802-077-9
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CURRENT FILING DATE: 2001-03-08
PRIOR APPLICATION NUMBER: US 08/405,617
PRIOR FILING DATE: 1995-03-15
PRIOR APPLICATION NUMBER: US 08/185,899
PRIOR FILING DATE: 1994-01-26
PRIOR PPLICATION NUMBER: PC7/US92/06860
PRIOR APPLICATION NUMBER: PC7/US92/06860
PRIOR FILING DATE: 1992-08-14
PRIOR PILING DATE: 1992-08-14
PRIOR APPLICATION NUMBER: US 07/879,495
PRIOR APPLICATION NUMBER: US 07/879,495
PRIOR APPLICATION NUMBER: US 07/744,768
PRIOR PILING DATE: 1992-05-07
PRIOR APPLICATION NUMBER: US 07/744,768
PRIOR FILING DATE: 1991-08-14
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Best Local &
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SEQ ID NO 9
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Best Local (
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APPLICANT: Presta, Leonard G.
TITLE OF INVENTION: Method of Treating Allergic Disorders
FILE REFERENCE: P0718P2C2US
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LOCATION: 1-114
OTHER INFORMATION: Light chain sequence derived
                                                                                                                                                                                                                                                                                                                                                                                    ORGANISM: Artificial sequence FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                TYPE: PRT
                                                                                                                                                                                                                                                                                                                                                                    OTHER INFORMATION: humanized maell, version 1, light chain
                                                                                                                                                                                                                                                                                                                                                                                                                                                LENGTH: 218
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                                                                                                                                          61 GVPSRFSGSGSGTDFTLTISSLQPEDFATYYCQHSWGIPRTFGQGTKVEIKRTV
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86.0%;
                                                                                                                                                                                                                                                                  81.8%; Score 486; DB 9;
86.0%; Pred. No. 1.6e-31;
tive 3; Mismatches 13
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Pred. No. 8.8e-32;
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61 GVPSRFSGSGSGTDFTLTISSLQPEDFATYYCQHSWGIPRTFGQGTKVEIKRTV

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PRIOR APPLICATION NUMBER: US 09/887,352
PRIOR FILING DATE: 1997-07-02
PRIOR APPLICATION NUMBER: US 08/887,352
PRIOR APPLICATION NUMBER: US 09/296,005
PRIOR PILING DATE: 1999-04-21
NUMBER OF SEQ ID NOS: 44
SEQ ID NO 13
LENGTH: 218
TYPE: PRT
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CURRENT FILING DATE: 2001-03-08
PRIOR APPLICATION NUMBER: US 08/405,617
PRIOR APPLICATION NUMBER: US 08/405,617
PRIOR APPLICATION NUMBER: US 08/185,899
PRIOR FILING DATE: 1994-01-26
PRIOR APPLICATION NUMBER: PCT/US92/06860
PRIOR FILING DATE: 1992-08-14
PRIOR APPLICATION NUMBER: US 07/879,495
PRIOR PILING DATE: 1992-05-07
PRIOR PILING DATE: 1992-05-07
PRIOR PILING DATE: 1991-08-14
NUMBER: US 07/744,768
PRIOR FILING DATE: 1991-08-14
NUMBER: OF SEQ ID NOS: 64
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                                                                                                         Query Match
Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                    TITLE OF INVENTION: Improved Anti-IgE Antibodies (as amended)
FILE REFERENCE: P1123CSUS
CURRENT APPLICATION NUMBER: US/09/920,171
CURRENT FILING DATE: 2001-08-01
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Lowman, Henry B. APPLICANT: Presta, Leonard
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ORGANISM: Artificial sequence
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PITLE OF INVENTION: Method of Preventing the Onset of Allergic Disorders
PILE REFERENCE: P0718P2C3US
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                                                                                                                                                                                          OTHER INFORMATION: Light chain sequence derived from MAE11
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98; Conserv
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DIQLTQSPSSLSASVGDRVTITCRASQSVDYDGDSYMNWYQQKPGKAPKLLIYAASYLES
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Jardieu, Paula M.
                                                                                     Conservative
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                                                                                                     Score 486; DB 9;
Pred. No. 1.6e-31;
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Pred. No. 1.6e-31
3; Mismatches 1
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PRIOR APPLICATION NUMBER: US 08/887,352
PRIOR PILING DATE: 1997-07-02
PRIOR PILING DATE: 1999-04-21
PRIOR PILING DATE: 1999-04-21
PRIOR APPLICATION NUMBER: US 09/920,171
PRIOR APPLICATION NUMBER: US 09/920,171
PRIOR PILING DATE: 2001-08-01
NUMBER OF SEQ ID NOS: 44
SEQ ID NO 13
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PRIOR FILING DATE: 1995-03-15
PRIOR PELICATION NUMBER: US 08/185,899
PRIOR FILING DATE: 1994-01-26
PRIOR APPLICATION NUMBER: PCT/US92/06860
PRIOR FILING DATE: 1992-08-14
PRIOR PELICATION NUMBER: US 07/879,495
PRIOR FILING DATE: 1992-05-07
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Best Local S
Matches 98
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PRIOR FILING DATE: 1991-08-14
NUMBER OF SEQ ID NOS: 68
SEQ ID NO 9
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Publication No.
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                                                                                                                                                                                          TITLE OF INVENTION: Improved Anti-IgE Antibodies FILE REFERENCE: F1123C3US CURRENT APPLICATION NUMBER: US/10/113,996 CURRENT FILLING DATE: 2002-04-01
                                                                                                                                                                                                                                                                                              APPLICANT:
                                                                                                                                                                                                                                                                                                                                         APPLICANT: Lowman, Henry B. APPLICANT: Presta, Leonard
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APPLICANT: Presta, Leonard G.
TITLE OF INVENTION: Anti-1GE Antibodies (as amended)
FILE REFERENCE: PO718P2CLDICIUS
CURRENT APPLICATION NUMBER: US/09/925,179
CURRENT FILING DATE: 2001-08-08
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Lowe, John
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86.0%;
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Pred. No. 1.6e-31;
3; Mismatches 13;
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; TYDE: PRT; ; CRANISM: Artificial Sequence; ; PEATURE: ; PEATURE: ; PEATURE: ; PEATURE: ; OTHER INFORMATION: Light chain sequence derived from MAE11

US-10-113-996-13

Query Match 81.8%; Score 486; DB 14; Length 218; Best Local Similarity 86.0%; pred. No. 1.6e-31; Matches 98; Conservative 3; Mismatches 13; Indels 0; Gaps 0; Matches 98; Conservative 3; Mismatches 13; Indels 0; Gaps 0; Matches 98; Conservative 3; Mismatches 11; Indels 0; Gaps 0; Matches 98; Conservative 3; Mismatches 13; Indels 0; Gaps 0; Matches 98; Conservative 3; Mismatches 11; Indels 0; Gaps 0; Matches 98; Conservative 3; Mismatches 13; Indels 0; Gaps 0; Matches 98; Conservative 3; Mismatches 13; Indels 0; Gaps 0; Matches 98; Conservative 3; Mismatches 13; Indels 0; Gaps 0; 0
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A Klausner R.D., Collins F.S., Wagner L., Shenmen C.M., Schuler G.D.,

A Altschul S.F., Zeeberg B., Butcow K.H., Schaefer C.F., Bhat N.K.,

A Altschul S.F., Zeeberg B., Moore T., Max S.I., Wang J., Hsieh F.,

A Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,

A Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,

A Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,

A Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,

A Stapleton M., Soares M.B., Bonaldo M.F., Casavant T.L., Schetz T.E.,

B Stapleton M., Joudin T.B., Toshiyuki S., Carninci P., Prange C.,

A Bosak S.A., McZwan P.J., McKernan K.J., Malek J.A., Gunaratne P.H.,

A Bosak S.A., McZwan P.J., McKernan K.J., Malek J.A., Gunaratne P.H.,

A Hilalon D.K., Muzny D.M., Sodergren E.J., Lu X., Gibbs R.A.,

A Villalon D.K., Muzny D.M., Sodergren E.J., Lu X., Gibbs R.A.,

A Wilting M., Madan A., Young A.C., Shevchenko Y., Bouffard G.G.,

A Whiting M., Madan A., Young A.C., Shevchenko Y., Bouffard G.G.,

A Whiting M., Touchman J.W., Green B.D., Dickson M.C.,

Bakesley R.W., Touchman J.W., Green B.D., Dickson M.C.,

A Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M., Sutterfield Y.S.,

A Tones S. J. Malek U., Smailus D.E., Schnerch A., Schein J.B.,

Jones S. J. Malek U., Smailus D.E., Schnerch A., Schein J.B.,
                                                                                                                                Query Match
Best Local S
Matches 89
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Best Local (
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAH18747;
02-MAR-2004 (TrEMBLrel.
02-MAR-2004 (TrEMBLrel.
02-MAR-2004 (TrEMBLrel.
                                                                                                                                                                                                                                                                                                                                                                                         TISSUE=Primary B-Cells;
Strausberg R.;
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Mammalia; Eutheria;
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                                                                                                                                                                                                                                                            Hypothetical protein. SEQUENCE 470 AA; 51715
                                                                                                                                                                                                                                                                                                                            EMBL; BC018747; AAH18747.1;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TISSUE=Primary B-Cells;
MEDLINE=22388257; PubMed=12477932;
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EVQLVESGGGLVQPGGSLRLSCVVSGFTFSSYWMSWVRQAPGKGLEWVANIKQDGSEKYY
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nilarity 72.4%;
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Primates;
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Pred. No. 8.3e-39;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AB092163; BAD00525.1;
  AF035024;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  60
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 61
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                           Immunol.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1 DVQLVESGGGLVQPGGSLRLSCAASGFTFSNYAISWVRQAPGKGLEWVSRINSD-GSTYY
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121 AA;
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(TrEMBLrel. 27, Last sequence update)
(TrEMBLel. 27, Last annotation update)
(In heavy chain VHDJ region (Fragment).
                                                                                                                                                                                                                                                                                                                                     (TrEMBLrel. 13, Created)
(TrEMBLrel. 13, Last sequence update)
(TrEMBLrel. 26, Last annotation updat
(Tremunoglobulin heavy chain variab
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                           Immunopathol.
     AAD56260.1;
                                                                                                                   PubMed=9614934;
n der Merwe P.L.,
                                                                                                                                                                                                                                           Chordata;
Primates;
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71.1%;
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                           87:184-192(1998)
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                                                                                                                                                                                                                                           Craniata; Vertebrata; Euteleostomi; Catarrhini; Hominidae; Homo.
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RESULT 5
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Best Local (
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Pfam; PF00047; ig; 1.
SMART; SM00406; IGv; 1.
PROSITE; PS50835; IG_LIKE; 1.
NON_TER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Honda T., Akahori Y., Kurosawa Y.;
"Libraries of heavy-chain antibodies
gamma3 in vivo repertoires.";
Submitted (SEP-2002) to the EMBL/GenE
EMBL; AB092052; BAD00414.1; -.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               BAD00414
BAD00414;
                     Q6MZV7;
05-JUL-2004 (TrEMBLrel. 27, Created)
05-JUL-2004 (TrEMBLrel. 27, Last sequence update)
05-JUL-2004 (TrEMBLrel. 27, Last annotation update)
Hypothetical protein DKFZp686C11235.
Name=DKFZp686C11235;
Homo sapiens (Human)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NON TER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TISSUE=Spleen;
Honda T., Akahori Y.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Eukaryota; Metazoa;
Mammalia; Eutheria;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               02-MAR-2004 (TrEMBLrel. 27, Created)
02-MAR-2004 (TrEMBLrel. 27, Last sequence update)
02-MAR-2004 (TrEMBLrel. 27, Last annotation update)
Immunoglobulin heavy chain VHDJ region (Fragment).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PIR; 8
HSSP;
                                                                                                                                                                                     Q6MZV7
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Camelus dromedarius
  Eukaryota; Metazoa; Chordata;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SEQÜENCE
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                                                                                                                                                                                                                                                                                                                                         62
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     13
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                             . Similarity
83; Conser
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                                                                                                                                                                                                                                                                                               DSVKGRFTISRDNAKNTLYLQMNSLKSGDTALYYCATSSSNSVDYWGQGTQVTVSSES
                                                                                                                                                                                                                                                                                                                           QRFKGRFTISLDKSKRTAYLQMNSLRAEDTAVYYCASWISDFFDYWGQGTLVTVSSAS
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113 AA;
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nilarity 70.3%;
Conservative 1:
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nilarity 73.5%;
Conservative
                                                                                                                                                                                   PRELIMINARY;
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12724 MW;
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Pred. No. 1.8e
11; Mismatches
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Pred. No. 1.
Craniata; Vertebrata;
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.8e-37;
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.7e-37;
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  Euteleostomi;
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InterPro; IPR003597; Ig-1ike.
InterPro; IPR003597; Ig-cl.
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InterPro; IPR003596; Ig_MC.
InterPro; IPR003596; Ig_V.
Pfam; PF00047; C1-set; 3.
Pfam; PF00047; IG; 2.
SMART; SM00409; IG; 2.
SMART; SM00407; IGC1; 3.
SMART; SM00407; IGC1; 3.
SMART; SM00407; IGC1; 3.
SMART; SM00409; IG LIKE; 4.
PROSITE; PS50835; IG_LIKE; 4.
PROSITE; PS50835; IG_LIKE; 4.
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02-MAR-2004 (TrEMBLrel. 27, Created)
02-MAR-2004 (TrEMBLrel. 27, Last sequence update)
02-MAR-2004 (TrEMBLrel. 27, Last annotation update)
Hypothetical protein DKFZp686C11235.
                                                                                                                                                                                                                                                                        TISSUE-Human small intestine;
Bloecker H., Boecher M., Mewes H.W., Weil B.,
Fobo G., Han M., Wiemann S.;
Submitted (AUG-2003) to the EMBL/GenBank/DDBJ
EMBL; BX640853; CAE45920.1; -.
Hypothetical protein.
SEQUENCE 473 AA; 52121 MW; 9476EAE4C0BFC44
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Submitted (AUG-2003) to the EMBL/GenBank/DDBJ databases
EMBL; BX640853; CAE45920.1; -.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Eukaryota; Metazoa;
Mammalia; Eutheria;
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SEQUENCE 473 AA; 5
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32; Conservative
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                                    EVQLVESGGGLVQPGGSLRLSCATSGYTFTEYIIHWVRQAPGKGLEWVASINPDYDITNY
EIQLVESGGGLVQPGGSLRLSCAASGFTFSSFEMNWVRQAPGKGLEWLSYITRSGNTVYY
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Boecher M., Mewes H.W.,
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Primata; C
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                                                                                                                                       13;
                                                                                                                                       Score 427.5;
Pred. No. 9.8e
13; Mismatches
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Pred. No. 9.8e-37;
.3; Mismatches 24;
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; Hominidae;
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RESULT 9
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TISSUE-Spleen;
TISSUE-TISSUE-TISSUE T., Akahori Y., Kurosawa Y.;
Honda T., Akahori Y., Kurosawa Y.;
"Libraries of heavy-chain antibodies re
"gamma3 in vivo repertoires.";
I gamma3 in vivo repertoires.";
L Submitted (SEP-2002) to the EMBL/GenBar
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02-MAR-2004
02-MAR-2004
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BAD00444;
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Mammalia; Eutheria;
NCBI_TaxID=9838;
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Submitted (SEP-2002) to the EMBL/GenF
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Mammalia; E
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02-MAR-2004 (TrEMBLrel. 27,
Immunoglobulin heavy chain V
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13856 MW;
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Chordata; Craniata; Vertebrata; Euteleostomi;
Cetartiodactyla; Tylopoda; Camelidae; Camelus
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Pred. No. 2.5e
14; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                             Last sequence update)
Last annotation update)
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Best Local S
Matches 84
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Q9UL71;
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SMART; SM00406; IGv; 1.
PROSITE; PS50835; IG_LIKE;
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HSSP; P01852; 1NFD
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                                                                                                                                                                                                                                                                                                                                                                                                                                               NCBI_TaxID=9606;
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Mammalia; Eutheria;
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01-OCT-2003 (TrEMBLrel. 25, Last annotation update)
Myosin-reactive immunoglobulin heavy chain variable
                                                                                                                                                                                                                SEQUENCE
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InterPro; IPR003596; Ig_v.
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121
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AF035043; AAD56279.1; -.
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121 AA;
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128 AA;
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Primates;
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66.7%;
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                                                                                                                                                                                                                13154 MW;
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                                                                                                                                                           9,
                                                                                                                                                          Score 426; DB 2
Pred. No. 3e-37;
9; Mismatches
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Pred. No. 2.5e-37;
2; Mismatches 27
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Craniata; Vertebrata; Euteleostomi; Catarrhini; Hominidae; Homo.
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Best Local S
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Matches 82
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02-MAR-2004
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Honda T., Akahori Y., Kurosawa Y.;
"Libraries of heavy-chain antibodies reflecting
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Cetartiodactyla; Tylopoda; Camelidae; Camelus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          02-MAR-2004 (TrEMBLrel. 27, Created)
02-MAR-2004 (TrEMBLrel. 27, Last sequence update)
02-MAR-2004 (TrEMBLrel. 27, Last annotation update)
02-MAR-2004 (TrEMBLrel. 27, Last annotation update
Immunoglobulin heavy chain VHDJ region (Fragment).
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Submitted (SEP-2002) to the I
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Cetartiodactyla; Tylopoda; Camelidae; Camelus
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NQRFKGRFTISLDKSKRTAYLQMNSLRAEDTAVYYCASWI---SDFFDYWGQGTLVTVSS
                                                                                                                                             EVQLVESGGGLVQPGGSLRLSCATSGYTFTEYIIHWVRQAPGKGLEWVASINPDYDITNY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SSES
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        EVQLVESGGGLVQPGGSLRLSCATSGYTFTEYIIHWVRQAPGKGLEWVASINPDYDITNY
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                                                                                                           EVQLVESGGGLVQPGGSLRLSCAASGFTFSKFWMHWVRQAPGKGLEWVSGINPVGINTYY
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(TrEMBLrel.
                                                                                                                                                                                                                                           Conservative
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67.7%;
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Last annotation update)
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Last annotation update)
                                                                                                                                                                                                                                    Score 425.5; |
Pred. No. 3.4e
|4; Mismatches
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Pred. No. 3e-3
LO; Mismatches
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AC Q8WUK
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WREDLINE-2238857; PubMedel2477932;

RA Strausberg R.L., Feingold E.A., Grouse L.H., Derge J.G.,

RA Klausner R.D., Collins F.S., Wagner L., Shenmen C.M., Schuler G.D.,

RA Altschul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,

RA Altschenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,

RA Diatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,

RA Stapleton M., Soares M.B., Bonaldo M.F., Casavant T.L., Scheetz T.E.,

RA Brownstein M.J., Uedin T.B., Toshiyuki S., Carninci P., Prange C.,

RA Raha S.S., Loquellano N.A., Peters G.J., Abramson R.D., Mullahy S.J.,

RA Roha S.S., Loquellano N.A., Peters G.J., Abramson R.D., Mullahy S.J.,

RA Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,

RA Villalon D.K., Muzny D.M., Sodergren E.J., Lu X., Gibbs R.A.,

RA Villalon D.K., Muzny D.M., Sodergren E.J., Lu X., Gibbs R.A.,

RA Pahey J., Helton B., Ketteman M., Madan A., Rodrigues S., Sanchez A.,

RA Mhiting M., Madan A., Young A.C., Shevchenko Y., Bouffard G.G.,

RA Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M., Butterfield Y.S.,

RA Krzywinski M.I., Skalska U., Smailus D.E., Schnerch A., Schein J.E.,

Tonas S. Marra M.A.,
                                                                                                                                                                                                                                                                                         Query Match
Best Local S
Matches 86
                                                                                                                                                                                                                                                                                                                                                      Strausberg R.;
Submitced (APR-2001) to the
EMBL; BC006402; AAH06402.1;
Hypothetical protein.
SEQUENCE 479 AA; 52281 MV
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAH06402;
02-MAR-2004 (TrEMBLrel.
02-MAR-2004 (TrEMBLrel.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Jones S.J., Marra M.A.; "Generation and initial analysis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TISSUE=Primary B-Cells;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SEQUENCE FROM N.A.
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Mammalia; Eutheria; Primates;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Hypothetical protein. Homo sapiens (Human).
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                                                                                                                                                             ADSVKGRFTISRDNAKKSLYLOMNSLRAEDTALYYCAKDIELDIVVVPAPMRGSLGFDYW
                                                                                                                                                                                           NORFKGRFTISLDKSKRTAYLOMNSLRAEDTAVYYCASWI-----
                                                                                                                                                                                                                             EVOLVESGGGLVQPGRSLRLSCAASGFTFDDYAMHWVRQVPGKGLEWVSGISCNSGSIGY
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                                                                                                                                                                                                                                                                                           Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SC1.
                                                                                                                                                                                                                                                                                                                                                         52281 MW;
                                                                                                                                                                                                                                                                                         67.1%; Score 425.5; DB 2 65.2%; Pred. No. 1.6e-36; bive 10; Mismatches 23
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27,
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Last annotation update)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Created)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Craniata; Vertebrata; Euteleostomi; Catarrhini; Hominidae; Homo.
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                                                                                                                                                                                           SDFFDYW
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Q8WUK1;

PRELIMINARY;

PRT;

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RX MEDLINE-22380257; PubMed=12477932;

RA Strausberg R.L., Feingold E.A., Grouse L.H., Derge J.G.,

RA Klausner R.D., Collins F.S., Wagner L., Shenmen C.M., Schuler G.D.,

RA Altschul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,

RA Hopkins R.F., Jordan H., Moore T., Max. S.I., Wang J., Hsieh F.,

RA Hopkins R.F., Jordan H., Moore T., Max. Rubin G.M., Hong L.,

RA Diatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,

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RA Raha S.S., KocEwan P.J., McKernan K.J., Malek J.A., Gunaratur P.H.,

RA Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,

Willalon D.K., Muzny D.M., Sodergren E.J., Lu X., Gibbs R.A.,

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RA Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M., Butterfield Y.S.,

RA Jones S.J., Marra M.A.;

"Generation and initial analysis of more than 15,000 full-length human

"Generation and initial analysis of more than 15,000 full-length human
                                                                                                                           Query Match
Best Local S
Matches 86
                                                                                                                                                                                                                                                                                                                                                                  EMBL; BCO2024

PIR; F36005;

PIR; G36005;

PIR; PH1642;

PIR; PH1643;

PIR; PH1646;

PIR; PH1646;

PIR; PH10098;

PIR; PL10098;

PIR; PL1020;

PIR; S15590;

PIR; S31116;

PIR; S31116;

PIR; S31116;

PIR; S31116;
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Homo sapiens (Human).
Eukaryota, Metazoa; Chordata,
Mammalia, Eutheria, Primates;
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Strausberg R.;
Submitted (DEC-2001) to the
                                                                                                                                                                                      PROSITE; PS50835; IG LIKE; 5.
PROSITE; PS00290; IG_MHC; UNKNOWN
SEQUENCE 613 AA; 67295 MW; 600
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Pfam;
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InterPro; IPR003597; Ig_C1.
InterPro; IPR00306; Ig_WHC.
InterPro; IPR003596; Ig_V.
Pfam; PF07654; C1-set; 4.
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PF00047; ig; 1.
; SM00406; IGv; 1.
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                                                                                                                                          Similarity
                                                          QVQLVESGGGVVQPGRSLRLSCAASGFTFSSYGMHWVRQAPGKGLEWVAVISYDGSNKYY
                NQRFKGRFTISLDKSKRTAYLQMNSLRAEDTAVYYCA-SWIS--DFFDYWGQGTLVTVSS
                                                                                 EVQLVESGGGLVQPGGSLRLSCATSGYTFTEYIIHWVRQAPGKGLEWVASINPDYDITNY
ADSVKGRFT1SRDNSKNTLYLQMNSLRAEDTAVYYCAKDWSEGVETFD1WGQGTMVTVSS
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(TrEMBLrel.
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G36005.
PH1642.
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PH1620.
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Pred. No. 2.8e-36;
3; Mismatches 25
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60C7F5950671E315 CRC64;
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Search completed: December 29, Job time: 133.79 secs
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Mammalia; Eutheria;
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                                                                                                                                                                                                                                                                                                            NCBI_TaxID=9838;
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                                              ADSVKGRFTISRDNAKEMVYLQMNSLKPEDTAVYYCVKWGDFSMDYWGEGTLVTISSES
                                                                                           EVOLVESGGGLVOPGGSLRLSCIASGFTFTGSWMWWVRQAPRKGLEWVSLINPSGSSTNY
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(TTEMBLrel. 27, Last sequence update)
(TTEMBLrel. 27, Last annotation update)
lin heavy chain VHDJ region (Fragment).
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Chordata; Craniata; Vertebrata; Euteleostomi;
Cetartiodactyla; Tylopoda; Camelidae; Camelus
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7; Mismatches
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      C;Keywords: nerc
F;15-98/Domain:
                         A;Cross-references: EMBL:X62955
A;Note: the nucleotide sequence was submitted to C;Superfamily: immunoglobulin V region; immunogloc;Keywords: heterotetramer; immunoglobulin
                                                                                  A;Status: preliminary; nucleic acid sequence
A;Molecule type: mRNA
A;Residues: 1-119 <RAA>
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65.5	65.5	65.7	65.7	65.7	65.8	65.8	65.9	65.9	66.2	66.4	66.4	66.5	66.6	66.6	66.6
151	122	141	122	118	160	121	122	127	121	121	120	114	134	117	147
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A60943	869910	S31669	S31675	S31105	S05271	S31104	S20772	S38489	<b>I55673</b>	S31118	S36273	S31120	S31679	S78486	I37780
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ALIGNMENTS

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### RESULT 2 S31107 Ig heavy chain - human C;Species: Homo sapiens (man) C;Species: Homo sapiens (man) C;Date: 02-Dec-1993 #sequence\_revision 26-May-1995 #text\_change 17-Mar-1999 C;Accession: S31107 R;Raaphoret, F.M.; Timmers, E.; Kenter, M.J.H.; van Tol, M.J.D.; Vossen, J.M.; Schuurma Eur. J. Immunol. 22, 247-251, 1992 Eur. J. Immunol. 22, 247-251, 1992 A;Title: Restricted utilization of germ-line V(H)3 genes and short diverse third comple A;Title: Restricted utilization of germ-line V(H)3 genes and short diverse third comple A;Reference number: S31104; MUID:92111633; PMID:1730252 A;Accession: S31107 A;Accession: S31107 S31598 1g heavy chain V region - human C;Species: Homo sapiens (man) C;Species: Homo sapiens (man) C;Date: 22-Nov-1993 #sequence\_revision 10-Nov-1995 #text\_change 23-Jul-1999 C;Accession: S31598 C;Accession: S31598 C;Accession: A.M.; Gauthier, L.; Boubli, L.; Pougereau, M.; Tonnelle, C. submitted to the EMBL Data Library, June 1992 submitted to the EMBL Data Library, June 1992 A;Cross-references: EMBL:Z14170; NID:g31001; PIDN:CAA78539.1; PID:g31002 C;Superfamily: immunoglobulin V region; immunoglobulin homology C;Keyvords: heterotetramer; immunoglobulin C;Keyvords: heterotetramer; immunoglobulin P;31-114/Domain: immunoglobulin homology <IMM> A; Reference number: S31585 A; Accession: S31598 A; Description: Mechanisms that generate human immunoglobulin diversity operate from 61 NQRFKGRPTISLDKSKRTAYLQMNSLRAEDTAVYYCA--SWISDFFDYWGQGTLVTVSS 117 1 EVQLVESGGGLVQPGGSLRLSCATSGYTFTEYIIHWVRQAPGKGLEWVASINPDYDITNY n 70.3%; Similarity 74.8%; ADSVKGRFTISRDNSKNTLYLQMNSLRAEDTAVYYCAKISWEVSRFDYWGQGTLVTVSS QVQLVESGGGVVQPGGSLRLSCAASGFTFSSYGMHWVRQAPGKGLEWVAFIRYDGSNKYY Conservative Score 446; DB 2; Pred. No. 2.2e-34; 5; Mismatches 23 23; Indels Length 135; 2 Gaps 76 60

homology

mitted to the EN immunoglobulin

EMBL Data in homology

Data Library,

October 1991

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Ig heavy chain V region (M43) - human C;Species: Homo sapiens (man) c;Date: 21-Dec-1990 #sequence_revision 21-Dec-1990 #text_change 16-Dec-1998 C;Accession: D36005 R;Schroeder Jr., H.W.; Wang, J.Y. Proc. Natl. Acad. Sci. U.S.A. 87, 6146-6150, 1990 Proferential utilization of conserved immunoglobulin heavy chain variable A;Reference number: A36005; MUID:90349571; PMID:2117273 A;Accession: D36005
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Proc. Natl. Acad. Sci. U.S.A. 87, 6146-6150, 1990
A;Title: Preferential utilization of conserved immunoglobulin heavy chain variable A;Reference number: A36005; MUID:90349571; PMID:2117273
A;Accession: C36005
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C;Keywords: heterotetramer; immunoglobulin
F;15-98/Domain: immunoglobulin homology <IMM>
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A;Molecule type: mRNA
A;Residues: 1-119 <SCH>
A;Cross_references: GB:M18513
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A;Map position: 14q32.33-14q32.33
C;Superfamily: immunoglobulin v region; immunoglobulin homology
C;Keywords: heterotetramer; immunoglobulin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     A;Status: preliminary
A;Molecule type: mRNA
A;Residues: 1-119 <SCH>
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                                                                                                                             Superfamily: immunoglobulin V region; immunoglobulin homology; Keywords: heterotetramer; immunoglobulin; 15-98/Domain: immunoglobulin homology <IMM>
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  EVQLVESGGGLVQPGGSLRLSCATSGYTFTEYIIHWVRQAPGKGLEWVASINPDYDITNY
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Pred. No. 1.1e-33;
9; Mismatches 22
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                                         Score 437; DB 2;
Pred. No. 1.3e-33;
7; Mismatches 21
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Pred. No. 4.5e-34;
7; Mismatches 22
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R,Mortari, F.; Wang, J.; Schroeder, H.W. submitted to the EMBL Data Library, April 1992 A;Description: Analysis of the IgA and IgG real A;Reference number: S20765 A;Accession: S20782
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A;Map position: 14G22.33-14G32.33
C;Superfamily: immunoglobulin V region; immunoglobulin C;Keywords: heterotetramer; immunoglobulin
                                                                                                                                                                                                                                                                                                                                                             R;Schroeder Jr., H.W.; Wang, J.Y.
Proc. Natl. Acad. Sci. U.S.A. 87, 6146-6150, 1990
A;Title: Preferential utilization of conserved immunoglobulin heavy
A;Reference number: A36005; MUID:90349571; PMID:2117273
                                                                                                                                                                                                                                                                                                                                                                                                                                                    Ig heavy chain V region (M74) - human C;Species: Homo sapiens (man) C;Date: 21-Dec-1990 #sequence_revision
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C;Species: Homo sapiens (man)
C;Date: 20-Feb-1995 #sequence_revision 20-Feb-1995
                                                                                                                                  C; Keywords: het
F; 15-98/Domain:
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A;Molecule type: mRNA
A;Residues: 1-121 <SCH>
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A; Residues: 1-124 < MOR>
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87; Conserv
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71.9%;
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69.4%;
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Pred. No. 1.5e
10; Mismatches
                                                    Score 436; Ub .,
Pred. No. 1.6e-33;
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Ig heavy chain - human

C; Species: Homo sapiens (man)

C; Species: Homo sapiens (man)

C; Species: 02-Dec-1993 #sequence_revision 26-May-1995 #text_change 17-Mar-1999

C; Accession: S31114

R; Raaphoret, F.M; Timmers, E.; Kenter, M.J.H.; van Tol, M.J.D.; Vossen, J.M.; Seur. J. Immunol. 22, 247-251, 1992

A; Title: Restricted utilization of germ-line V(H)3 genes and short diverse thire A; Reference number: S31104; MUID:92111633; PMID:1730252

A; Accession: S31114

A; Status: preliminary; nucleic acid sequence not shown; translation not shown A; Molecule type: mBNA

A; Molecule type: mBNA

A; Residues: 1-123 <RAA>

A; Cross-references: EMBL:X62963

A; Note: the nucleotide sequence was submitted to the EMBL Data Library, October C; Superfamily: immunoglobulin v region; immunoglobulin homology

C; Roywords: heterotetramer; immunoglobulin homology <IMM>
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S31686
                                                                                                                                                                                                                                                                                                                               Ig heavy chain V region - human (fragment)
C;Species: Homo sapiens (man)
C;Date: 22-Nov-1993 #sequence_revision 10-Nov-1995 #text_change 23-Jul-1999
C;Accession: S31686
C;Accession: S31686
R;Cuisinier, A.M.; Gauthier, L.; Boubli, L.; Fougereau, M.; Tonnelle, C.
                                                                                                                                               A; Molecule type: mRNA
A; Residues: 1-140 <CUI>
A; Residues: 1-140 <CUI>
A; Cross-references: EMBL: Z14205; NID: g30969; PIDN: CAA78574.1;
C; Superfamily: immunoglobulin V region; immunoglobulin homolog
C; Keywords: heterotetramer; immunoglobulin
C; Keywords: heterotetramer; immunoglobulin
                                                                                                                                                                                                                                             A;Description: Mechanisms that generate human immunoglobulin A;Reference number: S31585 A;Accession: S31686 A;Status: preliminary
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                                                            Matches
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Best Local
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                                                                              Similarity
EVQLVESGGGLVQPGGSLRLSCATSGYTFTEYIIHWVRQAPGKGLEWVASINPDYDITNY
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADSVKGRFTISRDNSKNTLYLQMNSLRAEDTAVYYCAKASLYLRFLEWL---FDYWGQGT 117
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NQRFKGRFTISLDKSKRTAYLQMNSLRAEDTAVYYCA-----SWISDFFDYWGQGT 111
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            EVQLVESGGGLVQPGGSLRLSCATSGYTFTEYIIHWVRQAPGKGLEWVASINPDYDITNY
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           QVQLVESGGGVVQPGRSLRLSCAASGFTFSSYAMHWVRQAPGKGLEWVAVISYDGSNKYY
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                                                                                                                                 immunoglobulin homology
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                                                                          68.8%;
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                                                          9;
                                                        Score 436; DB
Pred. No. 1.9e
9; Mismatches
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Pred. No. 1.7e-33;
8; Mismatches 19
                                                                                                                                                                    immunoglobulin homology
                                                                                                                                 <IMM>
                                                                                                                                                                                                                                                                                                                           1992
                                                                          DB 2;
..9e-33;
                                                        21;
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A;Status: preliminary
A;Molecule type: mRNA
A;Molecule type: mRNA
A;Residues: 1-132 <CUI>
A;Cross-references: EMBL:Z14168; NID:g30999; PIDN:CAA78537.1; PID:g31000
C;Superfamily: immunoglobulin V region; immunoglobulin homology
C;Keywords: heterotetramer; immunoglobulin
P;30-113/Domain: immunoglobulin homology <IMM>
                                                                                                                                                                                                                                                                                                                                                                                                                        Ig heavy chain V region - human
C;Species: Homo sapiens (man)
C;Date: 03-Mar-1994 #sequence_revision 10-Nov-1995 #text_change 23-Jul-1999
C;Accession: S31603
R;Cuisinier, A.M.; Gauthier, L.; Boubli, L.; Fougereau, M.; Tonnelle, C.
submitted to the EMBL Data Library, June 1992
A;Description: Mechanisms that generate human immunoglobulin diversity oper.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        A;Note: the nucleotide sequence was submitted to C;Superfamily: immunoglobulin V region; immunoglo C;Keywords: heterotetramer; immunoglobulin F;15-98/Domain: immunoglobulin homology <IVM>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    C;Species: Homo sapiens (man)
C;Date: 02-Dec-1993 #sequence_revision 26-May-1995 #text_change 17-Mar-1999
C;Accession: S31108
R;Raaphorst, F.M.; Timmers, B.; Kenter, M.J.H.; van Tol, M.J.D.; Vossen, J.P.
Eur. J. Immunol. 22, 247-251, 1992
A;Title: Restricted utilization of germ-line V(H)3 genes and short diverse tar. A;Reference number: S31104; MUID:92111633; PMID:1730252
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S31108
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A; Accession: S31603
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A;Status: preliminary; nucleic acid sequence
A;Molecule type: mENNA
A;Residues: 1-119 <RAA>
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Best Local S
Matches 86
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                                                                                                                                                      Similarity 72.6
85; Conservative
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NQRFKGRETISLDKSKRTAYLQMNSLRAEDTAVYYCASWISDFFDYWGQGTLVTVSS
                                                                  QVQLVESGGGVVQPGRSLRLSCAASGFTFSSYGMMVRQAPGKGLEWVAVISYDGSNKYY
                                                                                               EVQLVESGGGLVQPGGSLRLSCATSGYTFTEYIIHWVRQAPGKGLEWVASINPDYDITNY
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72.3%;
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Pred. No. 2e-3
9; Mismatches
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                                                                                                                                                    Score 435; DB 2;
Pred. No. 2.2e-33;
8; Mismatches 24
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RESULT 13
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                                                                                                                                                                                                                                                                                                                                                                                                         A; Molecule type: mRNA
A; Residues: 1-120 < MAH>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     R;Mahmoudi, M.; Edwards, J.; Cairns, E.; Bell, D. submitted to the EMBL Data Library, October 1994 A;Description: Molecular characterization of natural human A;Reference number: $48797
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Ig heavy chain V region (anti-Sm, VH3/Dxp4/JH4b) - C;Species: Homo sapiens (man) C;Date: 13-Jan-1995 #sequence_revision 13-Jan-1995 C;Accession: S48798
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A;Title: In vitro assembly of repertoires of antibody chains
A;Reference number: S46390; MUID:94254092; PMID:8196048
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A;Status: preliminary
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;Species: Homo sapiens (man)
;Date: 27-Jan-1995 #sequence_revision 27-Jan-1995 #text_change
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                                                                                                                                                           NQRFKGRFTISLDKSKRTAYLQMNSLRAEDTAVYYCAS----WISDFFDYWGQGTLVTVS 116
                                                                                                                                       ADSVKGRFTISRDNSKNTLYLOMNSLRAEDTAVYYCAKDRGFW-SGYKDYWGQGTLVTVS
                                                                                                                                                                                                                                        EVOLVESGGGLVQPGGSLRLSCATSGYTFTEYIIHWVRQAPGKGLEWVASINPDYDITNY
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Pred. No. 2.2e-33;
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Pred. No. 2.1e-33;
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RESULT 15
S70442
Ig heavy chain I
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F;15-98/Domain:
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A; Residues: 1-128 < MO
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                                                                                                                                                                                                                                                                                                                                                                                                                                                         A; Status: preliminary
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Best Local &
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   precursor V region (mu) -
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R;Mortari, F.; Newton, J.A.; Wang, J.Y.; Schroeder Jr., H.W. Bur. J. Immunol. 22, 241-245, 1992
A;Title: The human cord blood antibody repertoire. Frequent usage A;Reference number: S26786; MUID:92111632; PMID:1730251
A;Accession: S26786
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                A;Molecule type: mRNA
A;Residues: 1-140 <CUI>
A;Residues: 1-140 <CUI>
A;Cross-references: EMBL:Z14200; NID:g30957; PIDN:CAA78569.1; PID:g30958
C;Superfamily: immunoglobulin V region; immunoglobulin homology
C;Keywords: heterotetramer; immunoglobulin
F;34-117/Domain: immunoglobulin homology <IMM>
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C;Species: Homo sapiens (man)
C;Date: 22-Nov-1993 #sequence_revision 10-Nov-1995 #text_change 23-Jul-1999
C;Accession: S31588
C;Accession: S31588
R;Cuisinier, A.M.; Gauthier, L.; Boubli, L.; Fougereau, M.; Tonnelle, C.
submitted to the EMBL Data Library, June 1992
                                                                                                                                                                                                                                                                                                                                                                        A;Cross-references: EMBL:X61014; NID:g32800; PIDN:CAA43348.1; PID:g1335129 C;Superfamily: immunoglobulin V region; immunoglobulin homology C;Keywords: heterotetramer; immunoglobulin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    C;Species: Homo sapiens (man)
C;Date: 13-Jan-1995 #sequence_revision 13-Jan-1995 #text_change 20-Jun-2000
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                                                                                                                                                                            1 EVOLVESGGGLVQPGGSLRLSCATSGYTFTEYIIHWVRQAPGKGLEWVASINPDYDITNY
                                                   ADSVKGRFTISRDNAKNSLYLQMNSLRAEDTAVYYCARGLYCSSTSCYIWSNNWFDPWGQ
                                                                                               NQRFKGRFTISLDKSKRTAYLQMNSLRAEDTAVYYCAS------
                                                                                                                                                  QVQLVESGGGLVKPGGSLRLSCAASGFTFSDYYMSWIRQAPGKGLEWVSYISSSSSYTNY
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10; Mismatches
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Pred. No. 2
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fam

human (fragment)

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C:Species: Homo sapiens (man)
C:C:Date: 24-Jul-1998 #sequence_revision 24-Jul-1998 #text_change 09-Jul-2004
C:Accession: S70442
R;Cutesinier: A.M.; Funcux, F.; Fougereau, M.; Tonnelle, C.
Mol: Immunol. 29, 1363-1373, 1992
A;Title: IgM kappa/lambda EBV human B cell clone: an early step of differentiation of fell immunol. 29, 1363-1373, 1992
A;Fitle: IgM kappa/lambda EBV human B cell clone: an early step of differentiation of fell immunol. 29, 1363-1373, 1992
A;Fitle: IgM kappa/lambda EBV human B cell clone: an early step of differentiation of fell immunol. 29, 1363-1373, 1992
A;Fitle: IgM kappa/lambda EBV human B cell clone: an early step of differentiation of fell immunol. 29, 1363-1373, 1992
A;Fitle: IgM kappa/lambda EBV human B cell clone: an early step of differentiation of fell immunol. 29, 1992
A;Fitle: IgM kappa/lambda EBV human B cell clone: an early step of differentiation of fell immunol. 29, 1992
A;Fitle: IgM kappa/lambda EBV human B cell clone: an early step of differentiation of fell immunol. 29, 1992
A;Fitle: IgM kappa/lambda EBV human B cell clone: an early step of differentiation of fell immunol. 29, 2084VXI
A;Fitle: IgM kappa/lambda EBV human B cell clone: an early step of differentiation of fell immunol. 29, 2084VXI
C;Gucesion: S70442; MUID: 39024508; PMID:1383695
A;Fitle: IgM kappa/lambda EBV human B cell clone: Immunology cell immuno
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Result
No.
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Maximum I
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Maximum Match 100%
Listing first 45 summaries
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DB seq length: 2000000000
        Score
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1: /cgn2 6/ptodata/2/pubpaa/USO7_PUBCOMB.pep:*

2: /cgn2_6/ptodata/2/pubpaa/USO6_NEW_PUB.pep:*

3: /cgn2_6/ptodata/2/pubpaa/USO6_NEW_PUB.pep:*

4: /cgn2_6/ptodata/2/pubpaa/USO7_NEW_PUB.pep:*

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8: /cgn2_6/ptodata/2/pubpaa/USO8_NEW_PUB.pep:*
          Match Length
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/cgn2_6/ptodata/2/pubpaa/US10B_PUB.pep:*
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    14 US-01-044-896-5
15 US-09-811-384-10
15 US-10-404-286-10
15 US-10-404-286-10
16 US-09-940-166A-1
17 US-09-940-166A-1
18 US-09-940-166A-2
18 US-10-227-694-2
18 US-10-762-967-7
19 US-10-762-967-7
19 US-10-762-967-7
10 US-10-811-384-1
17 US-10-815-641-2
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17 US-10-818-765-4
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17	15	17	15	14	14	14	14	14	14	14	14	14	16	14	17	10	10	10	10	5	5	10	17	14	17	17	16	5	15	14	9
-10-723-003-5	-10-182-975-2	-10-723-003-5	-10-659-825-	-10-423-299-	-10-356-974-	-10-316-694-	-253-366-	US-10-364-953-14	-10-267-286A-1	-10-26	-10-227-	-10-020-786-	US-10-379-392-171		-10-835-641-	-09-795-798-	US-09-795-798-5	-09-726-258-7	-09-726-258-	-09-726-258	-09-726-258	-09-726-2	-10-835-643	0-423-299	-10-818-76	-10-719-31	0-619-754-	-10-600-152-	-10-608-626-	S-10-268-501	US-09-811-123-2
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## ALIGNMENTS

RESULT 1 US-10-044-896-5

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APPLICANT: Chuntharapai, Anan
APPLICANT: Kim, Jin K.
APPLICANT: Stewart, Timothy
APPLICANT: Presta, Leonard G.
TITLE OF INVENTION: ANTI-INTERFERON-ALPHA ANTIBODIES
FILE REFERENCE: GENERT.074A
CURRENT APPLICATION NUMBER: US/10/044,896
CURRENT FILING DATE: 2002-01-09
PRIOR APPLICATION NUMBER: 60/270775
PRIOR APPLICATION NUMBER: 60/270775
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                                                                                                                                                                                                                                             FEATURE:

10 OTHER INFORMATION: This sequence represents a humanized chimeric

10 OTHER INFORMATION: antibody comprising human and non-human sequence
                                                                                                                                                                                                                                                                                                                                                  NUMBER OF SEQ ID NOS: 14
SOPTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 5
LENGTH: 119
TYPE: PRT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 5, Application US/10044896 Publication No. US20030166228A1 GENERAL INFORMATION:
                                                                                                                                                  Query Match 100.0%; Score 634; DB 14; Best Local Similarity 100.0%; Pred. No. 1.8e-50; Matches 119; Conservative 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                              ORGANISM: Artificial Sequence
61 NQRFKGRFTISLDKSKRTAYLQMNSLRAEDTAVYYCASWISDFFDYWGQGTLVTVSSAS 119
                                                               non-human sequences.
                                                                                                                                                    Indels
                                                                                                                                                                                             Length 119;
                                                                                                                                                    <u>.</u>
                                                                                                                                                    Gaps
                                                                   60
                                                                                                          60
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RESULT 3
US-10-404-286-10
is Sequence 10, Application US/10404286
publication No. US20040057951A1
GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                        Matches
                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local &
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Patent No. US20020081294A1 GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        INFORMATION FOR SEQ ID NO: 10: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5 inch, 1.44 Mb flop;
MEDIUM TYPE: 3.5 inch, 1.44 Mb flop;
MEDIUM TYPE: 3.5 inch, 1.44 Mb flop;
COMPUTERS: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: WinPatin (Genentech)
CURRENT APPLICATION ADTA:
APPLICATION NUMBER: US/09/811,384
FILING DATE: 20-Dec-2000
CLASSIFICATION: CURROWN>
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICATION NUMBER: 09/251652
FILING DATE: 17-FEB-2000
APPLICATION NUMBER: 08/788800
FILING DATE: 22-JAN-1997
APPLICATION NUMBER: 60/093038
FILING DATE: 23-JAN-1996
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NAME: Love, Richard B.
REGISTRATION NUMBER: 34,659
REFERENCE/DOCKET NUMBER: P1729C1
TELECOMUNICATION INFORMATION:
TELEPHONE: 650/225-5530
TELEFAX: 650/952-9881
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Thomas, G. Roger
Gross, Cordell E.
TITLE OF INVENTION: ANTI-CD18 ANTIBODIES IN
NUMBER OF SEQUENCES: 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CORRESPONDENCE ADDRESS:
ADDRESSEE: Genente
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Bednar, Martin M.
                                                                                                                                                        121
                                                                                                                                                                                               114 TVSSAS 119
                                                                                                                                                                                                                                                                              19
                                                                                                                                                                                                                                       61 NQRFMDRFTISVDKSTSTAYMQMNSLRAEDTÄVYYCARWRGLNYGFDVRYFDVWGQGTLV 120
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97; Conserv
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                                                                                                                                                                                                                                                                                                                                                   EVQLVESGGGLVQPGGSLRLSCATSGYTFTEYIIHWVRQAPGKGLEWVASINPDYDITNY
                                                                                                                                                        TVSSAS 126
                                                                                                                                                                                                                                                                 NQRFKGRFTISLDKSKRTAYLQMNSLRAEDTAVYYCASWIS------DFFDYWGQGTLV
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CITY: South San Francisco
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADUKESSEE: Genentech, Inc. STREET: 1 DNA Way
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TOPOLOGY: Linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: Amino Acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NQRFKGRFTISLDKSKRTAYLQMNSLRAEDTAVYYCASWISDFFDYWGQGTLVTVSSAS 119
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                                                                                                                                                                                                                                                                                                                                                                                                    78.8%; ilarity 77.0%; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       232 amino acids
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                                                                                                                                                                                                                                                                                                                                                                                                    Score 499.5; DB 9
Pred. No. 7.3e-38;
7; Mismatches 15
                                                                                                                                                                                                                                                                                                                                                                                                                                             DB 9;
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                                                                                                                                                                                                                                                                                                                                                                                                                                             Length 232;
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                                                                                                         Sequence 1, Application US/09940166A Patent No. US20020058324A1
GENERAL INFORMATION:
APPLICANT: Blank, Gregory S.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
Best Local Similarity
Matches 97; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             INFORMATION FOR SEQ ID NO: 10: SEQUENCE CHARACTERISTICS: LENGTH: 232 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER READABLE FORM:

MEDIUM TYPE: 3.5 inch, 1.44 Mb floppy di
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOPTWARE: WinPatin (Genentech)
CURRENT APPLICATION NUMBER: US/10/404,286
FILING DATE: 31-Mar-2006
FILING DATE: 31-Mar-2006
CLASSIFICATION CAMPER: OS/611264
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICATION NUMBER: 09/811384
FILING DATE: 20-DEC-2000
APPLICATION NUMBER: 09/251652
FILING DATE: 17-FEB-2000
APPLICATION NUMBER: 08/788800
FILING DATE: 22-JAN-1997
APPLICATION NUMBER: 60/093038
FILING DATE: 23-JAN-1996
ATTORNEY, AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NAME: Evans, David W.
REGISTRATION NUMBER: NONE
REFERENCE/DOCKET NUMBER: P1729C2
TELECOMMUNICATION INFORMATION:
TELEPHONE: 650/225-1739
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Thomas, G. Roger
Gross, Cordell E.
TITLE OF INVENTION: ANTI-CD18 ANTIBODIES IN STROKE
NUMBER OF SEQUENCES: 15
                           Narindray, Daijit S.
Zapata, Gerardo A.
TITLE OF INVENTION: Protein Recovery
NUMBER OF SEQUENCES: 7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TOPOLOGY: Linear SEQUENCE DESCRIPTION: SEQ ID NO: 10:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Bednar, Martin
            CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CORRESPONDENCE ADDRESS:
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                                                                                                                                                                                                                                                                                                                       114 TVSSAS 119
                                                                                                                                                                                                                                                                                                                                                                                         61 NQRFKGRFTISLDKSKRTAYLQMNSLRAEDTAVYYCASWIS-----DFFDYWGQGTLV 113
                                                                                                                                                                                                                                                                                                                                                                61 NQRFMDRFTISVDKSTSTAYMQMNSLRAEDTAVYYCARWRGLNYGFDVRYFDVWGQGTLV
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1 EVQLVESGGGLVQPGGSLRLSCATSGYTFTEYIIHWVRQAPGKGLEWVASINPDYDITNY
                                                                                                                                                                                                                                                                            TVSSAS
                                                                                                                                                                                                                                                                                                                                                                                                                                               EVOLVESGGGLVOPGGSLRLSCATSGYTFTEYTMHWMRQAPGKGLEWVAGINPKNGGTSH
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TYPE: Amino Acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ADDRESSEE: Genentech, STREET: 1 DNA Way
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TELEFAX: 650/952-9881
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CITY: South San Francisco
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Conservative
                                                                                                                                                                                                                                                                              126
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77.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          7;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 499.5; DB 1
Pred. No. 7.3e-38;
7; Mismatches 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     DB 15;
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Indels Length 232;

7;

Gaps

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US-10-762-967-1
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                                                                                                                                                                                                                                                                                                                                                            Sequence 1, Application US/10762967 Publication No. US20040138426A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Matches 97; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TELEFAX: 650/952-9881
INFORMATION FOR SEQ ID NO: 1:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICATION NUMBER: 09/097,309
FILING DATE: 13-JUN-1997
ATTORNEY/AGENT INFORMATION:
NAME: Schwartz, Timothy R.
REGISTRATION NUMBER: 32171
REFERENCE/DOCKET NUMBER: P1105R1
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Match 78.8%;
Local Similarity 77.0%;
          COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5 inch, 1.44 Mb floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: WinPatin (Genentech)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5 inch, 1.44 Mb floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: WinPatin (Genentech)
CURRENT APPLICATION DATA:
                                                                                                                                                                                                              CORRESPONDENCE ADDRESS:
ADDRESSEE: Genentech, Inc.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICATION NUMBER: US/09/940,166A
FILING DATE: 27-Aug-2001
CLASSIFICATION: <Unknown>
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                             Zapata, Gerardo A. TITLE OF INVENTION: Protein Recovery
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SEQUENCE DESCRIPTION: SEQ ID NO: 1:
CURRENT
                                                                                                                                                                                                                                                                                                                   APPLICANT: Blank, Gregory S.
Narindray, Daljit S.
                                                                                                                                                                                                                                                            NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        121 TVSSAS 126
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            114 TVSSAS 119
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      61
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             61 NORFKGRFTISLDKSKRTAYLOMNSLRAEDTAVYYCASWIS------DFFDYWGQGTLV 113
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1 EVQLVESGGGLVQPGGSLRLSCATSGYTFTEYIIHWVRQAPGKGLEWVASINPDYDITNY 60
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NQRFMDRFTISVDKSTSTAYMQMNSLRAEDTAVYYCARWRGLNYGFDVRYFDVWGQGTLV 120
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 EVQLVESGGGLVQPGGSLRLSCATSGYTFTEYTMHWMRQAPGKGLEWVAGINPKNGGTSH 60
                                                                                                                                                        STATE: California
                                                                                                                                                                             STREET: 1 DNA Way
CITY: South San Francisco
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TYPE: Amino Acid
TOPOLOGY: Linear
                                                                                                                       ZIP: 94080
                                                                                                                                       COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   STATE: California
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  APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Score 499.5; DB 9;
Pred. No. 7.6e-38;
7; Mismatches 15;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   15;
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LENGTH: 241 amino acids

TYPE: Amino Acid

TOPOLOGY: Linear

SEQUENCE DESCRIPTION: SEQ ID NO: 1:
US-10-762-967-1
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US-09-940-166A-7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 7, Application US/09940166A Patent No. US20020058324A1 GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TELEFAX: 650/952-9881 INFORMATION FOR SEQ ID NO: 1:
                                                                                                                       ZIP: 94080

COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5 inch, 1.44 Mb floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: WinPatin (Genentech)
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/940,166A
FILING DATE: 27-Aug-2001
CLASSIFICATION: <UNKnown>
PRIOR APPLICATION DATA:
OLASSIFICATION DATA:
                                                               APPLICATION NUMBER: 09/097,309 FILING DATE: 13-JUN-1997 ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: Blank, Gregory S.

Narindray, Daljit S.
Zapata, Gerado A.
TITLE OF INVENTION: Protein Recovery
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NAME: Schwartz, Timothy R.
REGISTRATION NUMBER: 32171
REFERENCE/DOCKET NUMBER: P1105R1
TELECOMMUNICATION INFORMATION:
TELEPHONE: 650/225-7467
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/09/097,309
PILING DATE: 12-Jun-1998
APPLICATION NUMBER: 60/050951
PILING DATE: 13-JUN-1997
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                        CORRESPONDENCE ADDRESS:
ADDRESSEE: Genente
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    121 TVSSAS 126
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        61 NQRFMDRFTISVDKSTSTAYMQMNSLRAEDTAVYYCARWRGLNYGFDVRYFDVWGQGTLV 120
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                61
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                                                                                                                                                                                                                                                                                                                                                                  CITY: South San Francisco
STATE: California
COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           EVQLVESGGGLVQPGGSLRLSCATSGYTFTEYTMHWMRQAPGKGLEWVAGINPKNGGTSH 60
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FILING DATE: 21-Jan-2004
CLASSIFICATION: <Unknown>
NAME: Schwartz, Timothy R.
REGISTRATION NUMBER: 32171
REFERENCE/DOCKET NUMBER: P1105R1
                                                                                                                                                                                                                                                                                                                                                                                                                                      STREET: 1 DNA Way
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                        Genentech,
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77.0%;
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Pred. No. 7.6e-38;
7; Mismatches 15
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US-10-762-967-7
; Sequence 7, Application US/10762967
; Publication No. US20040138426A1
; GENERAL INFORMATION:
; APPLICANT: Blank, Gregory S.
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US-10-227-694-2
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                                                                                     RESULT 8
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ; SEQUENCE DESCRIPTION: SEQ ID NO: US-09-940-166A-7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PILE REFERENCE: p1867R1
CURRENT APPLICATION NUMBER: US/10/227,694
CURRENT FILING DATE: 2002-08-26
PRIOR APPLICATION NUMBER: US 60/315,209
PRIOR FILING DATE: 2001-08-27
NUMBER OF SEQ ID NOS: 6
SEQ ID NO 2
LENGTH: 300
TYPE: DET
                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 2, Application US/10227694
Publication No. US20030077739A1
GENERAL INFORMATION:
APPLICANT: Simmons, Laura
APPLICANT: Andersen, Dana
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match 78.8%;
Best Local Similarity 77.0%;
Matches 97; Conservative
                                                                                                                                                                                                                                                                                                                                      Matches
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TITLE OF INVENTION: A SYSTEM FOR ANTIBODY EXPRESSION AND ASSEMBLY
                                                                                                                                                                                                                                                                                                                                                                                                                                   LENGTH: 300
TYPE: PRT
ORGANISM: Artificial sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                  FEATURE:
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TELEPHONE: 650/225-7467
TELEFAX: 650/952-9881
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SEQUENCE CHARACTERISTICS:
                                                                                                                                      144
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                                                                                                                                   TVSSAS 149
                                                                                                                                                                                                    NQREMDRETISVDKSTSTAYMQMNSLRAEDTAVYYCARWRGLNYGFDVRYFDVWGQGTLV 143
                                                                                                                                                                                                                                                                   EVQLVESGGGLVQPGGSLRLSCATSGYTFTEYTMHWMRQAPGKGLEWVAGINPKNGGTSH
                                                                                                                                                                                                                                                                                              EVQLVBSGGGLVQPGGSLRLSCATSGYTFTEYIIHWVRQAPGKGLEWVASINPDYDITNY
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TYPE: Amino Acid
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                                                                                                                                                                                                                                                                                                                                  Conservative
                                                                                                                                                                                                                                                                                                                                            78.8%;
77.0%;
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Pred. No. 9.5e-38;
7; Mismatches 15;
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Pred. No. 9.5e-38;
7; Mismatches 15
                                                                                                                                                                                                                                                                                                                                                               DB 14; Length
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                                                                                                                                                                                                          Sequence 12, Application US/09811384
Patent No. US20020081294A1
GENERAL INFORMATION:
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Best Local S
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INFORMATION FOR SEQ ID NO: 7:
SEQUENCE CHARACTERISTICS:
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MEDIUM TYPE: 3.5 inch, 1.44 Mb floppy disk

COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: WinPatin (Genentech)
CURRENT APPLICATION NUMBER: US/10/762,967
PILING DATE: 21-Jan-2004
CLASSIFICATION INTROPER: US/10/762,967
PILING DATE: 21-Jan-2004
CLASSIFICATION NUMBER: US/09/097,309
PRIOR APPLICATION NUMBER: US/09/097,309
FILING DATE: 12-JUN-1998
APPLICATION NUMBER: 60/050951
PILING DATE: 13-JUN-1997
ATTORNEY/AGENT INFORMATION: D
                                                                      APPLICANT: Bednar, Martin M.
Thomas, G. Roger
Gross, Cordell B.
TITLE OF INVENTION: ANTI-CD18 ANTIBODIES
NUMBER OF SEQUENCES: 15
CORRESPONDENCE ADDRESS:
ADDRESSEE: Genentech, Inc.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NAME: Schwartz, Timothy R.
REGISTRATION NUMBER: 32171
REFERENCE/DOCKET NUMBER: P1105R1
TELECOMMUNICATION INFORMATION:
TELEPHONE: 650/225-7467
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Narindray, Daljit S.
Zapata, Gerardo A.
TITLE OF INVENTION: Protein Recovery
NUMBER OF SEQUENCES: 7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CORRESPONDENCE ADDRESS:
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                                                                                                                                                                                                                                                                                                                                                                                     114 TVSSAS 119
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                                                                                                                                                                                                                                                                                                                                                                                                                             84 NORFMORFTISVDKSTSTAYMOMNSLRAEDTAVYYCARWRGLNYGFDVRYFDVWGQGTLV
                                                                                                                                                                                                                                                                                                                                                                                                                                                   61 NQRFKGRFTISLDKSKRTAYLQMNSLRAEDTAVYYCASWIS------DFFDYWGQGTLV 113
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       24 EVOLVESGGGLVOPGGSLRLSCATSGYTFTEYTMHWMRQAPGKGLEWVAGINPKNGGTSH
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                                                        STREET: 1 DNA Way
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                    STATE: California
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        LENGTH: 300 amino acids
COUNTRY: USA
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                                    South San Francisco
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   78.8%;
77.0%;
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Pred. No. 9.5e-38;
7; Mismatches 15
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RESULT 10
US-10-404-286-12
; Sequence 12, Application US/10404286
; Publication No. US20040057951A1
; GENERAL INFORMATION:
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Best Local Similarity
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FILING DATE: 17-FEB-2000
APPLICATION NUMBER: 08/788800
FILING DATE: 22-JAN-1997
APPLICATION NUMBER: 60/093038
FILING DATE: 23-JAN-1996
ATTORNEY/ACENT INFORMATION:
COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5 inch, 1.44 Mb floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Windatin (Genentech)
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/404, 286
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SEQUENCE CHARACTERISTICS:
LENGTH: 450 amino acids
TYPE: Amino Acid
TOPOLOGY: Linear
SEQUENCE DESCRIPTION: SEQ ID NO:
                                                                                                                                                                                                                                                                                                   APPLICANT: Bednar, Martin M.
Thomas, G. Roger
Gross, Cordell E.
TITLE OF INVENTION: ANTI-CD18 ANTIBODIES IN STROKE
NUMBER OF SEQUENCES: 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NAME: Love, Richard B.
REGISTRATION NUMBER: 34,659
REFERENCE/DOCKET NUMBER: P1729C1
TELECOMMUNICATION INFORMATION:
TELEPHONE: 650/225-5530
                                                                                                                                                                                                                                                                                 CORRESPONDENCE ADDRESS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         114 TVSSAS 119
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   61 NQRFMDRFTISVDKSTSTAYMQMNSLRAEDTAVYYCARWRGLNYGFDVRYFDVWGQGTLV 120
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                                                                                                                                                                                             STREET: 1 DNA Way
CITY: South San Francisco
STATE: California
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        MEDIUM TYPE: 3.5 inch, 1.44 Mb 1
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOPTWARE: WinPatin (Genentech)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICATION NUMBER: US/09/811,384 FILING DATE: 20-Dec-2000 CLASSIFICATION: <Unknown>
                                                                                                                                                        ZIP: 94080
                                                                                                                                                                            COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TELEFAX: 650/952-9881
                                                                                                                                                                                                                                                         DDRESSEE: Genentech, Inc.
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77.0%;
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Pred. No. 1.4e-37;
7; Mismatches 15;
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US-10-835-641-23
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 23, Application US/10835641 Publication No. US20040236078A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Best Local Similarity 77.0
Matches 97; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NAME: EVANB, DAVID W
REGISTRATION NUMBER: NONE
REFRENCE/DOCKET NUMBER: P1
TELECOMMUNICATION INFORMATION:
TELEPHONE: 650/225-1739
TELEPAX: 650/925-1739
INFORMATION FOR SEQ ID NO: 12:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
CURRENT APPLICATION DATA:
APPLICATION UNMBER: US/10/835,641
FILING DATE: 30-Apr-2004
CLASSIFICATION: cUnknown>
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/09/705,398
FILING DATE: 02-Nov-2000
APPLICATION NUMBER: 08/146206
FILING DATE: 17-NOV-1993
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     FILING DATE: 31-Mar-2006
CLASSIFICATION: «Unknown»
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 09/811384
FILING DATE: 20-DEC-2000
APPLICATION NUMBER: 09/251652
FILING DATE: 17-EB-2000
APPLICATION NUMBER: 09/788800
APPLICATION NUMBER: 09/788800
FILING DATE: 22-JAN-1997
APPLICATION NUMBER: 60/93038
FILING DATE: 23-JAN-1996
ATTORNEY/AGENT INFORMATION:
NAME: RYANG DAYAG ME
                                                                                                                                                                                            COUNTRY: USA
ZIP: 94080
COMPUTER READABLE FORM:
COMPUTER: 3.5 inch, 1.44 Mb floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOPTWARE: WinPatin (Genentech)
                                                                                                                                                                                                                                                                                                                                                                                                                                    Presta, Leonard G.
TITLE OF INVENTION: Method for Making Humanized Antibodies
NUMBER OF SEQUENCES: 26
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TYPE: Amino Acid
TYPE: Amino Acid
TOPOLOGY: Linear
SEQUENCE DESCRIPTION: SEQ ID NO: 12:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: Carter, Paul J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     121 TVSSAS 126
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            114 TVSSAS 119
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 61 NORFKGRFTISLDKSKRTAYLOMNSLRAEDTAVYYCASWIS-----DFFDYWGQGTLV 113
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1 EVQLVESGGGLVQPGGSLRLSCATSGYTFTEYIIHWVRQAPGKGLEWVASINPDYDITNY 60
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               EVOLVESGGGLVQPGGSLRLSCATSGYTFTEYTMHWMRQAPGKGLEWVAGINPKNGGTSH 60
                                                                                                                                                                                                                                                                                                                                                        STREET: 1 DNA Way
CITY: South San Francisco
STATE: California
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77.0%;
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Pred. No. 1.4e-37;
7; Mismatches 15
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; LENGTH: 119
; TYPE: PRT
; ORGANISM: Murine
US-10-044-896-2
                   RESULT 13
US-10-818-765-4
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Sequence 4, Application US/10818765
                                                                                                                                                                                                                                                                   Query Match
Best Local S
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Publication No. US20030165228A1
GENERAL INFORMATION:
APPLICANT: Chuntharapai, Anan
APPLICANT: Kim, Jin K.
                                                                                                                                                                                                                                                                                                                                                                                                  SOFTWARE: FastSEQ for Windows Version 4.0 SEQ ID NO 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Best Local Similarity Matches 97; Conserv
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Best Local
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APPLICANT: Stewart, Timothy
APPLICANT: Presta, Leonard G.
TITLE OF INVENTION: ANTI-INTERPERON-ALPHA ANTIBODIES
FILE REFERENCE: GENENT.074A
CURRENT APPLICATION NUMBER: US/10/044,896
CURRENT FILING DATE: 2002-01-09
PRIOR APPLICATION NUMBER: 60/270775
PRIOR FILING DATE: 2001-02-22
NUMBER OF SEQ ID NOS: 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TELEFAX: 650/952-9881
INFORMATION FOR SEQ ID NO: 23:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                     Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICATION NUMBER: 07/715272
FILING DATE: 14-UN-1991
ATTORNEY/AGENT INFORMATION:
NAME: Lee, Wendy M.
REGISTRATION NUMBER: 40,378
REFERENCE/DOCKET NUMBER: P0709P1D2
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SEQUENCE DESCRIPTION: SEQ ID NO:
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                                                                                                                               61 NORFKGRETISLDKSKRTAYLOMNSLRAEDTAVYYCASWISDFFDYWGOGTLVTVSSAS 119
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                                                                                          NQRFKGKATLTLDKSSRTAYLELRSLTSEDSAVYYCASWISDFFDYWGQGTTLMVSAAS 119
                                                                                                                                                                     EVQLQQSGPELVKPGASVKISCKTSGYTFTEYIIHWVKQGHGRSLEWIGSINPDYDITNY
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       EVQLVESGGGLVQPGGSLRLSCATSGYTFTEYTMHWMRQAPGKGLEWVAGINPKNGGTSH 79
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TYPE: Amino Acid TOPOLOGY: Linear
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                                                                                                                                                                                                                                                     Conservative
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                                                                                                                                                                                                                                                               77.8%; Score 493; DB 14; Length 119; 73.1%; Pred. No. 1.4e-37;
                                                                                                                                                                                                                                                 17; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            7; Mismatches
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Pred. No. 1.5e-37;
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CURRENT APPLICATION NUMBER: US/09/811,123
CURRENT FILING DATE: 2001-03-16
PRIOR APPLICATION NUMBER: 60/238,327
PRIOR FILING DATE: 2000-10-05
PRIOR APPLICATION NUMBER: 09/602,530
PRIOR FILING DATE: 2000-06-23
NUMBER OF SEQ ID NOS: 11
SOPTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 2
LENGTH: 119
TYPE: PRT
ORGANISW: Artificial Sequence
FEATURE:
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                                                                                                                                                                                                      ; OTHER INFORMATION: Humanized Antibody Sequence US-09-811-123-2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ; OTHER INFORMATION: humanized sequence US-10-818-765-4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               GENERAL INFORMATION:
APPLICANT: Sharon Erickson
APPLICANT: Ralph Schwall
APPLICANT: Mark Sliwkowski
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 2, Application US/09811123 Patent No. US20020001587A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Publication No. US20040202658A1 GENERAL INFORMATION:
                                                                                                                                 Matches
                                                                                                                                                Query Match
Best Local (
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LENGTH: 452
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TITLE OF INVENTION: THERAPY OF AUTOIMMUNE DISEASE IN A PATIENT WITH TITLE OF INVENTION: INADEQUATE RESPONSE TO A TWF-A INHIBITOR FILE REFERENCE: P2027R1-US
CURRENT APPLICATION NUMBER: US/10/818,765
CURRENT FILING DATE: 2004-04-06
PRIOR APPLICATION NUMBER: US 60/461,4819
PRIOR PILING DATE: 2003-04-09
NUMBER OF SEQ ID NOS: 4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TITLE OF INVENTION: METHODS OF TREATMENT USING ANTI-EXB
TITLE OF INVENTION: ANTIBODY-MAYTANSINOID CONJUGATES
FILE REFERENCE: GENENT.073A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TYPE: PRT ORGANISM: Artificial sequence FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     118 VTVSSAS 124
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61 NQRFKGRFTISLDKSKRTAYLQMNSLRAEDTAVYYCASWI--SDFFDYWGQGTLVTVSS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        61 NQKFKGRPTISVDKSKNTLYLQMNSLRAEDTAVYYCARVVYYSNSYW----YFDVWGQGTL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               61 NQRFKGRFTISLDKSKRTAYLQMNSLRAEDTAVYYCAS------WISDFFDYWGQGTL
                                                      93;
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97; Conserv
                                                                                                                                                    Similarity
                                                                                                                               Conservative
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                                                                                                                                                76.8%;
78.2%;
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76.4%;
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                                                                                                                               Score 487; DB 9;
Pred. No. 5.1e-37;
1; Mismatches 13
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Pred. No. 6.3e-37;
6; Mismatches 13;
                                                                                                                                   13;
                                                                                                                                                                   Length 119
                                                                                                                                 Indels
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                                                                                                                               Gaps
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RESULT 15

US-10-268-501-4

Sequence 4, Application US/10268501

Publication No. US/2003006924A1

GENERAL INFORMATION:

APPLICANTON SILWANDER: US/10/268,501

CURRENT FILING DATE: 2002-10-10

PRIOR PPLICATION NUMBER: US/10/268,501

CURRENT FILING DATE: 2002-10-10

PRIOR PPLICATION NUMBER: US/10/268,501

CURRENT FILING DATE: 2002-10-10

PRIOR PPLICATION NUMBER: US/10/268,501

CURRENT FILING DATE: 2002-06-23

PRIOR PPLICATION NUMBER: US/10/268,501

CURRENT FILING DATE: 1000-06-23

PRIOR PILING DATE: 1000-06-23

PRIOR PILING DATE: 1000-06-23

PRIOR PILING DATE: 1099-06-25

PRIOR PILING DATE: 2002-06-23

PRIOR PILING
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sequence

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Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      seq length: 0
seq length: 2000000000
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Gapop 10.0 ,
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634
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275.930 Million cell updates/sec
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1: /cgn2_6/ptodata/1/1aa/5A_COMB.pep:*
2: /cgn2_6/ptodata/1/1aa/5B_COMB.pep:*
3: /cgn2_6/ptodata/1/1aa/6B_COMB.pep:*
4: /cgn2_6/ptodata/1/1aa/6B_COMB.pep:*
5: /cgn2_6/ptodata/1/1aa/PCTUS_COMB.pep:*
6: /cgn2_6/ptodata/1/1aa/backfIles1.pep:*
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US-07-934-373C-30
US-08-437-642B-30
PCT-US93-07832-30
PCT-US93-07832-30
PCT-US93-07832-31
US-09-097-309-1
US-09-097-309-1
US-09-097-171A-1
US-09-097-171A-1
US-09-097-171A-1
US-09-097-171A-1
US-09-097-171A-1
US-09-097-171A-1
US-09-097-171A-1
US-09-097-171A-1
US-09-097-171A-3
US-09-460-587-7
US-09-460-587-7
US-09-460-587-7
US-09-460-587-7
US-09-460-587-3
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US-08-437-642B-23
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US-09-705-392A-23
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Sequence 27, Appl
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Sequence 23, Appli
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                                                                                                                                                    COMPUTER READABLE FORM:

MEDIUM TYPE: 3.5 inch, 1.44 Mb floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Winderin (Genentech)
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/07/934,373C
FILING DATE: 21-Aug-1992
CLASSIFICATION NUMBER: PCT/US92/05126
PRIOR APPLICATION NUMBER: PCT/US92/05126
PRIOR APPLICATION NUMBER: PCT/US92/05126
PRIOR APPLICATION NUMBER: PCT/US92/05126
PRIOR APPLICATION NUMBER: 07/115272
PRIOR APPLICATION NUMBER: 07/115272
PRIOR APPLICATION NUMBER: 07/15272
PRIOR APPLICATION NUMBER: 07
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US-07-934-373C-27
                                       Query Match
Best Local Similarity
Matches 98; Conserva
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: Paul J. Carter
APPLICANT: Leonard G. Presta
TITLE OF INVENTION: Immunoglobulin Variants
NUMBER OF SEQUENCES: 48
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                STREET: 1 DNA Way
CITY: South San Francisco
STATE: California
COUNTRY: USA
ZIP: 94080
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US-07-934-373C-26
US-08-437-642B-26
PCT-US-93-07832-38
US-07-934-373C-36
US-07-934-373C-36
US-07-934-373C-36
US-07-934-373C-36
US-08-437-642B-35
US-08-437-642B-33
US-08-437-642B-33
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                                                  Score 503.5; DB 2;
Pred. No. 4.4e-42;
6; Mismatches 15;
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US-08-437-642B-27
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                                                                                                                  Matches
                                                                                                                                               Query Match
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MEDIUM TYPE: 3.5 inch, 1.44 Mb floppy di
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: WinPatin (Genentech)
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/437,642B
FILING DATE: 09-May-1995
CLASSIFICATION: 530
CLASSIFICATION: 530
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                GENERAL INFORMATION:
APPLICANT: Paul J. Carter
APPLICANT: Leonard G. Presta
TITLE OF INVENTION: Immunoglobulin Variants
NUMBER OF SEQUENCES: 47
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                TELEFAX: 650/952-9881
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                          APPLICATION NUMBER: 07/7:
FILING DATE: 14-JUN-1991
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICATION NUMBER: PCT/US92/05126
FILING DATE: 15-JUN-1992
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PRIOR APPLICATION DATA:
APPLICATION NUMBER:
FILING DATE: 17-NOV-
                                                                                                                                                                                                                                               SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                            TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/9:
FILING DATE: 21-AUG-1992
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                         NAME: Lee, Wendy M.
REGISTRATION NUMBER: 40,378
REFERENCE/DOCKET NUMBER: PO
                                                                                                                                 Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COUNTRY: U
ZIP: 94080
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NQRFKGRFTISLDKSKRTAYLQMNSLRAEDTAVYYCASWIS------DFFDYWGQGTLV 113
                                                                EVQLVESGGGLVQPGGSLRLSCATSGYTFTEYIIHWVRQAPGKGLEWVASINPDVDITNY 60
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                                                EVQLVESGGGLVQPGGSLRLSCATSGYTFTEYTMHWMRQAPGKGLEWVAGINPKNGGTSH
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                                                                                                                            79.4%;
77.8%;
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                                                                                                               Score 503.5; DB 3
Pred. No. 4.4e-42;
6; Mismatches 15
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PCT-US93-07832-27
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CURRENT APPLICATION DATA:
APPLICATION NUMBER: PCT/US93/07:
FILING DATE: 1930820
CLASSIFICATION DATA:
APPLICATION DATA:
APPLICATION UMBER: 07/715272
APPLICATION UMBER: 07/715272
APPLICATION DATA:
APPLICATION DATA:
APPLICATION DATA:
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APPLICATION DATA:
APPLICATION DATA:
07/934373
APPLICATION DATA:
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TELEX: 910/371-7168
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CORRESPONDENCE ADDRESS:
ADDRESSEE: Genentech,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TITLE OF INVENTION: NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ATTORNEY/AGENT INFORMATION:
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REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER: IBM PC OPERATING SYSTEM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          MEDIUM TYPE:
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CITY: South San Francisco
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121 TVSSAS 126
                                     114 TVSSAS 119
                                                                                                       61 NQRFKGRFTISLDKSKRTAYLQMNSLRAEDTAVYYCASWIS------DFFDYWGQGTLV 113
                                                                             61 NQRFMDRFTISVDKSKNTLYLQMNSLRAEDTAVYYCARWRGLNYGFDVRYFDVWGQGTLV
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                                                                                                                                                                                   1 EVQLVESGGGLVQPGGSLRLSCATSGYTFTEYIIHWVRQAPGKGLEWVASINPDYDITNY 60
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                                                                                                                                                           1 EVQLVESGGGLVQPGGSLRLSCATSGYTFTEYTMHWMRQAPGKGLEWVAGINPKNGGTSH
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98; Conservative
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                                                                                                                                                                                                                                                                                                                                                   linear
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SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          5.25 inch,
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77.8%;
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Pred. No. 4.4e-42;
6; Mismatches 15;
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RESULT 4 US-07-934-373C-29

Sequence 29, Application US/07934373C Patent No. 5821337

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                                                Sequence 10, Application US/08788800
Patent No. 5914112
GENERAL INFORMATION:
APPLICANT: Bednar, Martin M.
APPLICANT: Thomas, G. Roger
APPLICANT: Grose, Cordell E.
TITLE OF INVENTION: ANTI-CD18 ANTIBODIES IN STROKE
NUMBER OF SEQUENCES: 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Matches
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APPLICANT: Paul J. Carter
APPLICANT: Leonard G. Presta
TITLE OF INVENTION: Immunoglobulin Variants
NUMBER OF SEQUENCES: 48
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TELEFAX: 650/952-9881
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
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PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/715272
FILING DATE: 14-JUN-1991
ATTORNEY/AGENT INFORMATION:
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COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5 inch, 1.44 Mb floppy disk
COMPUTER: IBM PC compatible
COMPUTER: PC-DOS/MS-DOS
COMPATTING SYSTEM: PC-DOS/MS-DOS
CORRESPONDENCE ADDRESS:
ADDRESSEE: Genentech, Inc.
STREET: 460 Point San Bruno Blvd
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      REGISTRATION NUMBER: 40,378
REFERENCE/DOCKET NUMBER: 0
TELECOMMUNICATION INFORMATION:
TELEPHONE: 650/225-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PRIOR APPLICATION DATA:
APPLICATION NUMBER: PCT/US92/05126
FILING DATE: 15-JUN-1992
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            LENGTH: 232 amino acids
TYPE: Amino Acid
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICATION NUMBER: US/07/934,373C
FILING DATE: 21-Aug-1992
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CITY: South San Francisco
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COUNTRY:
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                                                                                                                                                                                                                                                                                   TVSSAS 126
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HER: P0709P2
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US-08-788-800-10
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Patent No. 6054297
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TELEFAX: 415/952-9881
TELES: 910/371-7168
INFORMATION FOR SEQ ID NO: 10
SEQUENCE CHARACTERISTICS:
LENGTH: 232 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
Best Local
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                                                                                                            COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5 inch, 1.44 Mb floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: WinPatin (Genentech)
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Paul J. Carter
APPLICANT: Leonard G. Presta
TITLE OF INVENTION: Immunoglobulin Variants
NUMBER OF SEQUENCES: 47
                                 APPLICATION NUMBER: US/08/437,642B FILING DATE: 09-May-1995 CLASSIFICATION: 530 PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                     CORRESPONDENCE ADDRESS:
ADDRESSEE: Genentech
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NAME: Lee, Wendy M.
REGISTRATION NUMBER: 40,378
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MEDIUM TYPE: 3.5 inch, 1.44 Mb floppy disk
COMPUTER: IBM PC compatible
COMPUTER: COMPUTER: COMPATIBLE COMPUTER: COMPUTER: COMPUTER: COMPUTER: COMPUTER: COMPUTER: COMPUTER: COMPUTER: COMPUTER: COMP
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                                                                                                                                                                                                                                                                                               CITY: South San Francisco
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APPLICATION NUMBER: FILING DATE: 21-AU
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STATE: California
COUNTRY: USA
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97; Conservative
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77.0%;
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Pred. No. 1.1e-41;
7; Mismatches 15;
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PRIOR APPLICATION DATA:
APPLICATION NUMBER:
FILING DATE: 17-NOV-

08/146206

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RESULT 7
PCT-US93-07832-29
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INFORMATION FOR SEQ ID NO: 29
SEQUENCE CHARACTERISTICS:
LENGTH: 232 amino acids
TYPE: Amino Acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 29, Applications
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
                                                                                                                                                                                                       COMPUTER READABLE FORM:
MEDIUM TYPE: 5.25 inch, 360 Kb floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: patin (Genentech)
CURRENT APPLICATION NUMBER: PCT/US93/07832
APPLICATION NUMBER: PCT/US93/07832
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FILING DATE: 15-JUN-1992
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/715272
FILING DATE: 14-JUN-1991
ATTORNEY/AGENT INFORMATION:
NAME: Lee, Wendy M.
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                                                    PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/715272
APPLICATION NUMBER: 07/715272
PRIOR APPLICATION DATA:
APPLICATION NUMBER: PCT/US92/05126
PILING DATE: 15-JUN-1992
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TITLE OF INVENTION:
NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                         CORRESPONDENCE ADDRESS:
                                       PRIOR APPLICATION DATA:
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REGISTRATION NUMBER: 40,378
REFERENCE/DOCKET NUMBER: PO
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STREET: 460 Point San Bruno Blvd
CITY: South San Francisco
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APPLICATION NUMBER: 07/934373
FILING DATE: 21-AUG-1992
                                                                                                                                                                        CLASSIFICATION:
                                                                                                                                                                                           FILING DATE:
                                                                                                                                                                                                                                                                                                                                                             COUNTRY: USA
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l Similarity 77.0%;
97; Conservative
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Pred. No. 1.1e-41;
7; Mismatches 15;
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RESULT 8
US-07-934-373C-30
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Best Local Similarity
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TELEX: 910/371-7168
INFORMATION FOR SEQ ID NO: 3
SEQUENCE CHARACTERISTICS:
LENGTH: 232 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          GENERAL INFORMATION:
                                                                      FILING DATE: 14-JUN-1991
ATTORNEY/AGENT INFORMATION:
NAME: Lee, Wendy M.
REGISTRATION NUMBER: 40,378
REFERENCE/DOCKET NUMBER: P070
TELECOMMUNICATION INFORMATION:
TELEPHONE: 650/225-1994
                                    TELEFAX: 650/952-9881
INFORMATION FOR SEQ ID NO: 30:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    REGISTRATION NUMBER: 70: REFERENCE/DOCKET NUMBER: 70: TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Paul J. Carter
APPLICANT: Leonard G. Presta
TITLE OF INVENTION: Immunoglobulin Variants
NUMBER OF SEQUENCES: 48
CORRESPONDENCE ADDRESS:
ADDRESSE: Genentech, Inc.
                                                                                                                                                                                                                                                                                                                                                   COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5 inch, 1.44 Mb floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Winpatin (Genentech)
CURRENT APPLICATION DATA:
SEQUENCE CHARACTERISTICS
LENGTH: 235 amino aci
                                                                                                                                                                                                                      APPLICATION NUMBER: PCT/US92/05126
FILING DATE: 15-JUN-1992
PRIOR APPLICATION DATA:
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                                                                                                                                                                                                                                                                                             APPLICATION NUMBER: US/07/934,373C FILING DATE: 21-Aug-1992 CLASSIFICATION: 530
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                                                                                                                                                                                    APPLICATION NUMBER: 07/715272 FILING DATE: 14-JUN-1991
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77.0%;
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US-08-437-642B-30
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                                                     US-08-437-642B-30
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Matches 97, Conservative
                   Query Match
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 Best Local Similarity
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APPLICATION NUMBER: 07/934373
FILING DATE: 21-AUG-1992
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/146206
FILING DATE: 17-NOV-1993
                                                                                                                     INFORMATION FOR SEQ ID NO: 30: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Paul J. Carter
APPLICANT: Leonard G. Presta
TITLE OF INVENTION: Immunoglobulin Variants
NUMBER OF SEQUENCES: 47
CORRESPONDENCE ADDRESS:
                                                                                                                                                                     REFERENCE/DOCKET NUMBER: PO
TELECOMMUNICATION INFORMATION:
TELEPHONE: 650/225-1994
                                                                                                                                                                                                                                                     APPLICATION NUMBER: 07/7:
FILING DATE: 14-JUN-1991
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                            PRIOR APPLICATION DATA:
APPLICATION NUMBER: PCT/US92/05126
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/437,642B
FILING DATE: 09-May-1995
                                                                                                                                                                                                                                                                                                         PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     MEDIUM TYPE: 3.5 inc
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER: IBM PC
OPERATING SYSTEM:
                                                                                                                                                                                                                     NAME: Lee, Wendy M.
REGISTRATION NUMBER: 40,378
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CLASSIFICATION:
                                                                        TOPOLOGY:
                                                                                                       LENGTH:
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South San Francisco
California
                                                                                   amino Acid
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                                                                                                                                                        650/952-9881
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Pred. No. 1.1e-41;
7; Mismatches 15
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Score 499.5; DB 3; Pred. No. 1.1e-41;
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                Length 235;
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PCT-US93-07832-30
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                                                                                                                                  Matches
                                                                                                                                                               Query Match
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TELEX: 910/371-7168
INFORMATION FOR SEQ ID NO:
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APPLICATION NUMBER: 07/7
FILING DATE: 14-JUN-1991
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-
SOFTWARE: patin (Genentech)
CURRENT APPLICATION DATA:
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MEDIUM TYPE: 5.25 in
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                                                                                                                                                                                                                                                                                                                                                   REFERENCE/DOCKET NUMBER: 70
TELECOMMUNICATION INFORMATION
                                                                                                                                                                                                                                                                                                                                                                                                                     ATTORNEY/AGENT INFORMATION:
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CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                   LENGTH:
TYPE: a
                                                                                                                                               Local
                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICATION NUMBER: 07/934373
FILING DATE: 21-AUG-1992
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NORFMORFTISVDKSTSTAYMOMNSLRAEDTAVYYCARWRGLNYGFDVRYFDVWGQGTLV
                             NORFKGRFTISLDKSKRTAYLOMNSLRAEDTAVYYCASWIS-----DFFDYWGQGTLV 113
                                                                                    EVQLVESGGGLVQPGGSLRLSCATSGYTFTEYIIHWVRQAPGKGLEWVASINPDYDITNY 60
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                                                                EVOLVESGGGLVQPGGSLRLSCATSGYTFTEYTMHWMRQAPGKGLEWVAGINPKNGGTSH
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                                                                                                                                                 Score 499.5; DB 5
Pred. No. 1.1e-41;
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                                                                                                                                  Mismatches
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Best Local S
Matches 97
Sequence 1, Application US/09097171A
Patent No. 6171586
GENERAL INFORMATION:
APPLICANT: Lam, Xanthe M.
APPLICANT: Oeswein, James Q.
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APPLICATION NUMBER: 60/050951
PILING DATE: 13-UUM-1997
ATTORNEY/AGENT INFORMATION:
NAME: Schwartz, Tinothy R.
REGISTRATION NUMBER: 32171
REFERENCE/DOCKET NUMBER: 91105R
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        GENERAL INFORMATION:
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MEDIUM TYPE: 3.5 inch, 1.44 Mb floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: WinPatin (Genentech)
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/097,309
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SEQUENCE CHARACTERISTICS:
LENGTH: 241 amino acids
TYPE: Amino Acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: Blank, Gregory S.
APPLICANT: Narindray, Daljit S.
APPLICANT: Zapata, Gerardo A.
TITLE OF INVENTION: Protein Recovery
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CLASSIFICATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       12-Jun-1998
                                                                                                                                                                                                                                                                                                                                                                                                    78.8%;
77.0%;
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P1105R1
                                                                                                                                                                                                                                                                                                                                                                                     7; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                    Score 499.5; DB 3
Pred. No. 1.1e-41;
                                                                                                                                                                                                                                                                                                                                                                                                                       DB 3;
                                                                                                                                                                                                                                                                                                                                                                                                                     Length 241;
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Matches
                                                                                                                                                                                                     Sequence 1, Application US/09460587 Patent No. 6322997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
                                                                                                                                                                                     GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TELEFAX: 650/952-9881
INFORMATION FOR SEQ ID NO: 1:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/874
FILING DATE: 13-JUN-1997
ATTORNEY/AGENT INFORMATION:
NAME: Lee, Wendy M.
REGISTRATION NUMBER: 40,37
REFERENCE/DOCKET NUMBER: E
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT:
APPLICANT:
APPLICANT:
                                                                                                                 APPLICANT: Blank, Gregory S.
APPLICANT: Narindray, Daljit S.
APPLICANT: Zapata, Gerardo A.
TITLE OF INVENTION: Protein Recovery
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SEQUENCE CHARACTERISTICS:
LENGTH: 241 amino acids
TYPE: Amino Acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TELECOMMUNICATION INFORMATION:
TELEPHONE: 650/225-1994
TELEFAX: 650/952-9881
                                                                                NUMBER OF SEQUENCES: 7
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SOFTWARE: WinPatin (Ge:
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER READABLE FORM:
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CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TITLE OF
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADDRESSEE: Genentech, Inc.
STREET: 1 DNA Way
CITY: South San Francisco
                                                STREET:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    MEDIUM TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COUNTRY:
                                                                ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TOPOLOGY: Linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CLASSIFICATION:
                                                                                                                                                                                                                                                                                                            121 TVSSAS 126
                                                                                                                                                                                                                                                                                                                                           114 TVSSAS 119
                                                                                                                                                                                                                                                                                                                                                                            61 NQRFMDRFTISVDKSTSTAYMQMNSLRAEDTAVYYCARWRGLNYGFDVRYFDVWGQGTLV 120
                                                                                                                                                                                                                                                                                                                                                                                                    61 NORFKGRFTISLDKSKRTAYLOMNSLRAEDTAVYYCASWIS------DFFDYWGQGTLV
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    97;
                                                                                                                                                                                                                                                                                                                                                                                                                                              1 EVQLVESGGGLVQPGGSLRLSCATSGYTFTEYTMHMMRQAPGKGLEMVAGINPKNGGTSH
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              South San Fi
: California
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T: Weissburg, Robert P.
F: Wong, Rita L.
INVENTION: Antibody Formulation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      California
                                                1 DNA Way
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USA
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Shahrokh, Zahra
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                                                              Genentech, Inc.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    3.5 inch, 1.44 Mb floppy disk
                            Francisco
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                78.8%;
77.0%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 499.5; DB 3;
Pred. No. 1.1e-41;
7; Mismatches 15;
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RESULT 14
US-09-940-166A-1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 1, Application US/09940166A Patent No. 6716598
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ATTORNEY AGENT INFORMATION:
NAME: Schwartz, Timothy R.
REGISTRATION NUMBER: 32171
REFERENCE/DOCKET NUMBER: P1105R1
TELECOMMUNICATION INFORMATION:
TELEPHONE: 650/225-7467
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TELEFAX: 650/952-9881
INFORMATION FOR SEQ ID NO: 1:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SEQUENCE CHARACTERISTICS:
LENGTH: 241 amino acids
TYPE: Amino Acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DO
SOFTWARE: WinPatin (Genentech)
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5 inch, 1.44 Mb floppy disk
COMPUTER: IBM PC compatible
COMPUTER: PC DOS/MS-DOS
OPERATING SYSTEM: PC DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PRIOR APPLICATION DATA:
APPLICATION NUMBER:
                                          COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5 inch, 1.44 Mb floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARB: WinPatin (Genentech)
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/940,166A
PILING DATE: 27-Aug-2001
CLASSIFICATION - CUNKNOWN>
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Local Similarity 77.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TOPOLOGY: Linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CLASSIFICATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Blank, Gregory S.

Narindray, Daljit S.
Zapata, Gerardo A.
TITLE OF INVENTION: Protein Recovery
NUMBER OF SEQUENCES: 7
                                                                                                                                                                                                                                                                                                                                                                                                         CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           121 TVSSAS 126
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   114 TVSSAS 119
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                                                                                                                                                                                                                                                                                                                  CITY: South San Francisco
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NORFMORFTISVOKSTSTAYMOMNSLRAEDTAVYYCARWRGLNYGFDVRYFDVWGQGTLV 120
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                                                                                                                                                                                                                                                                                                COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                               STREET: 1 DNA Way
  APPLICATION NUMBER: 09/097,309 FILING DATE: 13-JUN-1997
                                                                                                                                                                                                                                                                           ZIP: 94080
                                                                                                                                                                                                                                                                                                                                                                                    ADDRESSEE: Genentech, Inc.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 499.5; DB 3;
Pred. No. 1.1e-41;
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                                                                                                                       COMPUTER READABLE FORM:

MEDIUM TYPE: 3.5 inch, 1.44 Mb flc
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: WinPatin (Genentech)
CURRENT APPLICATION DATA:
OPERATION NUMBER: US/09/097,309
FILING DATE: 12-Unn-1998
CLASSIFICATION NUMBER: 60/050951
APPLICATION NUMBER: 109/7
APPLICATION NUMBER: 109/7
ATTORNEY/ACENT INFORMATION:
NAME: Schwartz, Timothy R.
REGISTRATION NUMBER: 321/71
REFERENCE/DOCKET NUMBER: 3110SR1
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
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                                                               TELEFAX: 650/952-9881
INFORMATION FOR SEQ ID NO: 7:
SEQUENCE CHARACTERISTICS:
LENGTH: 300 amino acids
TYPE: Amino Acid
TOPOLOGY: Linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ZIP: 94080
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US/09/097,309

1.44 Mb floppy disk

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RESULT 15
US-09-097-309-7
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                                                                                                                                                                                                                                               Sequence 7, Application US/09097309
Patent No. 6121428
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Best Local Similarity 77.0 Matches 97; Conservative
                                                                                                                      GENERAL INFORMATION:
APPLICANT: Blank, Gregory S.
APPLICANT: Narindray, Daljit S.
APPLICANT: Zapata, Gerardo A.
TITLE OF INVENTION: Protein Rec
NUMBER OF SEQUENCES: 7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      INFORMATION FOR SEQ ID NO: 1:
SEQUENCE CHARACTERISTICS:
                                                                                                         CORRESPONDENCE ADDRESS:
                     STATE: California
                                             CITY: South San Francisco
                                                                STREET:
COUNTRY:
                                                                                    ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NAME: Schwartz, Timothy R.
REGISTRATION NUMBER: 32171
REFERENCE/DOCKET NUMBER: P1105R1
TELECOMMUNICATION INFORMATION:
TELEPHONE: 650/225-7467
TELEFAX: 650/952-9881
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SEQUENCE DESCRIPTION: SEQ ID NO: 1:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                               121 TVSSAS 126
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      61 NQRFKGRFTISLDKSKRTAYLQMNSLRAEDTAVYYCASWIS------DFFDYWGQGTLV 113
                                                                                                                                                                                                                                                                                                                                                                                                                                                              61 NQRFMDRFTISVDKSTSTAYMQMNSLRAEDTAVYYCARWRGLNYGFDVRYFDVWGQGTLV
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1 EVQLVESGGGLVQPGGSLRLSCATSGYTFTEYTMHWMRQAPGKGLEWVAGINPKNGGTSH 60
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         LENGTH: 241 amino acids
TYPE: Amino Acid
TOPOLOGY: Linear
                                                                  1 DNA Way
    USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  78.8%;
77.0%;
                                                                                                                                                  Protein Recovery
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Score 499.5; DB 4
Pred. No. 1.1e-41;
7; Mismatches 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       DB 4;
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Maximum Match 100%
Listing first 45 summaries
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Maximum DB
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                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence:
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                                                                                                                                                                                                                                                     seq length: 0
seq length: 2000000000
A_Geneseq_23Sep04:*
1: geneseqp1980s:*
2: geneseqp1990s:*
3: geneseqp2000s:*
4: geneseqp2001s:*
5: geneseqp2013as:*
6: geneseqp2013as:*
7: geneseqp2003bs:*
8: geneseqp2004s:*
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634
1 EVQLVESGGGLVQP
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                                                                                                                                                                                                                                                                                                                                                                                                                                 EVQLVESGGGLVQPGGSLRL.....ISDFFDYWGQGTLVTVSSAS 119
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

## SUMMARIES

25	24	23	22	21	20	19	18	17	16	15	14	13	12	11	10	9	8	7	σ	U	4	ω	N	۳	NO.	Result
487	487	487	487	487	487	493	499.5	499.5	499.5	499.5	499.5	499.5	499.5	499.5	499.5	•	499.5	499.5	499.5	499.5	499.5	499.5	499.5	634	Score	,
76.8	76.8	76.8	76.8	76.8	76.8	77.8	78.8	78.8	78.8	78.8	78.8	78.8	78.8	78.8	78.8	78.8	78.8	78.8	78.8	78.8	78.8	78.8	78.8	100.0	Match	Query
119	119	119	119	119	119	119				450	450	450	300	300	300	300	277	241	241	232	232	232	232	119	Match Length	
8	æ	σ	4.	4.	4	ហ	N	w	æ	S	N	N	σ	4	N	N	N	4	N	8	σ	N	N	σ		
ADJ88007	ADE71453	AAU74540	AAB61584	AAB60399	AAB62086	AAE28147	AAR30775	AAB03664	ADK18343	ABG31890	AAY08755	AAW34505	ABP72746	AAB66785	AAW30635	AAW95623	AAW30631	AAB66776	AAW95614	ADK18341	ABG31888	AAY08753	AAW34503	AAE28150	ID	
_	Ade71453 Humanise	Aau74540 Human ant	Aab61584 Humanised	Aab60399 Humanised	Aab62086 Humanised	Aae28147 Murine 9F		Aab03664 Anti-CD18	Amino		Aay08755 Human IgG			Aab66785 Protein e	Aaw30635 Recombina	Aaw95623 pS1130 ex	Aaw30631 Recombina	Aab66776 rhuMAb CD	Aaw95614 Humanized	Adk18341 Amino aci	Abg31888 Humanised	Aay08753 Human ant	Aaw34503 Heavy cha	nie	Description	

45	44	43	42	41	40	39	38	37	36	35	34	33	32	31	30	29	28	27	26
477.5	477.5	477.5	477.5	477.5	477.5	477.5	477.5	477.5	477.5	477.5	477.5	477.5	477.5	477.5	477.5	480.5	481	487	487
75.3	75.3	75.3	75.3	75.3	75.3	75.3	75.3	75.3	75.3	75.3	75.3	75.3	75.3	75.3	75.3	75.8	75.9	76.8	76.8
256	256	256	256	253	253	253	253	253	253	253	253	253	253	253	253	122	451	119	119
ω	w	N	N	7	7	σ	6	σ	σ	w	ω	w	w	N	N	N	œ	œ	œ
AAB30321	AAY77765	AAY29457	AAW69304	AAE39086	AAE39089	ABU59504	ABU59506	ABU13793	ABU13791	AAB30314	AAB30316	AAY77759	AAY77757	AAY29454	AAY29444	AAR30772	ADF11670	ADP43327	ADN12053
Aab30321	Aay77765	Aay29457	Aaw69304	Aae39086	Aae39089	Abu59504	Abu59506	Abu13793	Abu13791	Aab30314	Aab30316	Aay77759	Aay77757	Aay29454	Aay29444	Aar30772	Adf11670	Adp43327	Adn12053
Humanised	Humanised	6G4V11N35	Humanised	6G4-2-5V1	Humanised	Humanised	Humanised	Humanised	huxCD3v9,	anti-CD11	Humanised	Variable							

# ALIGNMENTS

AAE28150 standard; protein; 119 ₹

AAE28150;

27-DEC-2002 (first entry)

Humanised 9F3 version 13 (V13) heavy chain variable domain.

RESULT 1
AAE28100
ID AAE282
XX AAE28
XX AAE28
XX AAE20
XX Huma
XX Huma Human; interferon alpha; IFNalpha; insulin-dependent diabetes mellitus; autoimmune disease; systemic lupus erythematosus; autoimmune thyroiditis; antibody therapy; immunosuppressive; antiinflammatory; dermatological; 9F3 monoclonal antibody; heavy chain variable domain.

Homo sapiens. Synthetic.

×			Ϋ́				
		Region		Region		Region	Key
	<pre>/note= "Complementarity determining region 3 (CDR3)"</pre>	99106	<pre>/note= "Complementarity determining region 2 (CDR2)"</pre>	5066	<pre>/note= "Complementarity determining region 1 (CDR1)"</pre>	2635	Location/Qualifiers
	determining		determining		determining		
	region 3		region 2		region 1		
	(CDR3) "		(CDR2) "		(CDR1)"		

WO200266649-A2

29-AUG-2002.

29-JAN-2002; 2002WO-US002709.

22-FEB-2001; 2001US-0270775P. 09-JAN-2002; 2002US-00044896.

(GETH ) GENENTECH INC.

Chuntarapai A, Kim JK, Presta LG, Stewart T;

WPI; 2002-682767/73.

New anti-interferon alpha monoclonal antibody, 9F3, useful for treating autoimmune diseases e.g. insulin-dependent diabetes mellitus, systemic lupus erythematosus and autoimmune thyroiditis.

Disclosure; Page 98; 100pp; English.

The present invention relates to novel anti-interferon alpha (IFNalpha)

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RESULT 2
AAW34503
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Matches
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This sequence represents the heavy chain of the humanised H52 antibody. The H52 antibody can be used in the method of the invention. The method is for treating focal ischaemic stroke, i.e. damage to the brain due to interrupted blood supply, in a mammal, caused by obstruction of a main cerebral artery, comprises administering anti-CD18 antibody to increase cerebral blood flow and/or reduce the size of the area of brain that
                                                                                                                                                                                                                                               Administering an anti-CD18 antibody to treat focal ischaemic stroke - increasing cerebral blood flow and reducing size of brain infarct.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Humanised antibody; HuH52; heavy chain; focal ischaemic stroke; embolism; brain damage; anti-CD18 antibody; cerebral blood flow; thromboembolism; transient ischaemic attack; thrombolytic therapy; thrombosis; therapy; systemic hypoperfusion; cardiac arrest.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          of humanised H52 antibody.
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No. 4.1e-52;
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F δ 밁 S

arterial obstruction.

CD18

This invention describes a method for improving the clinical outcome in focal ischaemic stroke by administering novel anti-CD18 antibody which

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Col 29-32;

25pp; English

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                                                                                                                                                                                                                                                                                                                                                            IGG4; Kappa-CL domain; lambda-CL domain; focal ischaemic stroke;
cerebroprotective; cerebral artery obstruction; blood flow; infarct;
CD18 extracellular domain; endothelium; CD11b/CD18 complex dissociation;
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                                                                                                                              Thomas
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                                                                                                                                                                                                                                                                                                                                             antibody; huH52; heavy
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77.0%;
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No. 4.2e-39;
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RESULT 4
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22-JAN-1997;
17-FEB-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             focal ischaemic stroke; main cerebral artery;
tissue plasminogen activator; anti-CD18 antibody; stroke;
acute ischaemic stroke; thrombolytic therapy;
Fabv1b salvage receptor binding epitope; thromboembolic stroke
                     The invention relates to a method or reducing infarct size in focal
                                                                                           Claim
                                                                                                                                     Increasing cerebral blood flow and/or reducing infarct size in ischemic stroke using anti-CD18 antibody and tissue plasminogen is useful to improve clinical outcome in acute ischemic stroke.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human; humanisec focal ischaemic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Humanised
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ABG31888
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       27-JUN-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  US2002081294-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            05-NOV-2002
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Homo
                                                                                                                                                                                                                                                                                                                               (GETH )
    ø
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              humanised H52 antibody; huH52;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    97; Conserv
                                                                                                                                                                                                                                                                                                                               GENENTECH
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NORFMORFTISVDKSTSTAYMQMNSLRAEDTAVYYCARWRGLNYGFDVRYFDVWGQGTLV
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     EVQLVESGGGLVQPGGSLRLSCATSGYTFTEYIIHWVRQAPGKGLEWVASINPDYDITNY
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            H52
    cerebral
                                                                                                                                                                                                                                                                                 Gross
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      2000US-00811384.
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                                                                                                                                                                                                                                                                                                                                                                        96US-0093038P.
97US-00788800.
99US-00251652.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            antibody (huH52)
                                                                                           5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       126
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 119
                                                                                       27pp;
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    artery
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   peptide;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          78.8%;
77.0%;
                                                                                                                                                                                                                                                                                 Thomas
                                                                                           English
    ä
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      232
    ø
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 499.5; DB 2
Pred. No. 4.2e-39;
7; Mismatches 15
                                                                                                                                                                                                                                                                                 g,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Fab
                        of increasing cerebral ischaemic stroke caused
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       fragment, heavy
                                                                                                                                                                                                                                                                                 Gross
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              cerebral
comprising
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  co-administering
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                          caused
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            chain.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              flow;
                        blood flow and/
d by obstruction
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  232;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              infarct
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                                                                                                                                                               activator
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              size;
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Best Local
                                                                                                                                                                                23-JAN-1996;
22-JAN-1997;
17-FEB-1999;
20-DEC-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        plasminogen activator and anti-CD18 antibody about 3-5 hours af stroke. The method is used to improve the clinical outcome in a ischaemic stroke and to provide an alternative to thrombolytic for treating thromboembolic stroke, particularly where thrombol therapy has been unsuccessful or is contra-indicated. The presengence represents a humanised H52 antibody (huH52) Fab, heavy used in the method of the invention
                                                                                                                                                                                                                                  31-MAR-2003;
                                                                                                                                                                                                                                                                          US2004057951-A1
                                                                                                                                                                                                                                                                                                          Homo sapiens
                                                                                                                                                                                                                                                                                                                                      cerebral artery; tissue humanised H52 antibody;
                                                                                                                                                                                                                                                                                                                                                          Cerebral blood flow;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 232
                                                                                                                                          Bednar
                                                                                                                                                              (GETH )
                                                                                                                                                                                                                                                       25-MAR-2004.
                                                                                                                                                                                                                                                                                              Synthetic.
                                                                                                                                                                                                                                                                                                                                                                              Amino acid sequence
                                                                                                                                                                                                                                                                                                                                                                                                    17-JUN-2004
                                                                                                                                                                                                                                                                                                                                                                                                                       ADK18341;
                                                                                                                                                                                                                                                                                                                                                                                                                                          ADK18341 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             114
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                                                                                                                                                                                                                                                                                                                            heavy
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                                                                                                                                                              GENENTECH
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NORFMORFTISVDKSTSTAYMOMNSLRAEDTAVYYCARWRGLNYGFDVRYFDVWGQGTLV
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NORFKGRFTISLDKSKRTAYLOMNSLRAEDTAVYYCASWIS-----DFFDYWGQGTLV
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TVSSAS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TVSSAS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       EVQLVESGGGLVQPGGSLRLSCATSGYTFTEYIIHWVRQAPGKGLEWVASINPDYDITNY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          EVQLVESGGGLVQPGGSLRLSCATSGYTFTEYTMHWMRQAPGKGLEWVAGINPKNGGTSH
                                                                                                                                          Gross
                                                                                                                                                                                96US-0093038P.
97US-00788800.
99US-00251652.
2000US-00811384.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Conservative
                                                                                                                                                                                                                                   2003US-00404286
                                                                                                                                                                                                                                                                                                                             chain.
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                                                                                                                                                                                                                                                                                                                                                                                                   (first
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           126
                                                                                                                                                             INC
                                                                                                                                         œ,
                                                                                                                                                                                                                                                                                                                                                                                                                                         protein;
                                                                                                                                                                                                                                                                                                                                                                              of humanised H52 (huH52) Fab heavy chain
                                                                                                                                                                                                                                                                                                                                                                                                   entry)
                                                                                                                                                                                                                                                                                                                                                          infarct size; focal ischaemic stroke;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          78.8%;
77.0%;
                                                                                                                                          Thomas
                                                                                                                                                                                                                                                                                                                                      plasminogen activator; tPA; anti-CD18
huH52 Fab; stroke; cerebroprotective;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 499.5; | Pred. No. 4.2e 7; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                          232
                                                                                                                                         g,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           499.5; DB 5;
No. 4.2e-39;
                                                                                                                                          Gross
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   15;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            thrombolytic
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heavy chain
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                                                                                                                                                                                                                                                                                                                                      vasotropic;
                                                                                                                                                                                                                                                                                                                                                antibody;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                acute
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     therapy
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    120
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flow and/or reducing infarct size in focal ischaemic stroke caused by obstruction of a main cerebral artery in a mammal, particularly humans. The method comprises co-administering tiasue placemine actions actions.

co-administering

tissue plasminogen

Claim 11;

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present

invention

relates

to a method English

for

increasing

cerebral

plood

Increasing cerebral blood flow and/or reducing infarct size in ischemic stroke caused by obstruction of a main cerebral artery human comprises co-administering tissue plasminogen activator as

n focal ry in a and ant

2004-257111/24

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RESULT 6
AAW95614
AID 9AW9
AC AAW9
AC AAW9
AC AW9
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
Best Local &
   New stable aqueous antibody subjected to lyophilisation,
                                                                                                                                                                                                                                                                                                                                           13-JUN-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                12-JUN-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   17-DEC-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WO9856418-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Mus musculus.
                                                                                                        WPI; 1999-080860/07
                                                                                                                                                                       Weissburg
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Monoclonal antibody; heavy chain; humanized; myocardial infarction; burns; thermal injury; ischemic; shock; ischaemic; haemorrhagic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Humanized anti-CD18 antibody rhuMAb CD18 heavy chain.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            08-JUN-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAW95614;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  nemorrhagic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAW95614 standard; protein; 241
                                                                                                                                                                                                                                                                            (GETH ) GENENTECH INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 121
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97; Conser
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NQREMDRFTISUDKSTSTAYMQMNSLRAEDTAVYYCARWRGLNYGFDVRYFDVWGQGTLV
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TVSSAS 119
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            EVQLVESGGGLVQPGSSLRLSCATSGYTFTEYTWHWMRQAPGKGLEWVAGINPKNGGTSH
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         stroke.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Å
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                                                                                                                                                                   Wong
                                                                                                                                                                                                                                                                                                                                              97US-00874897
                                                                                                                                                                                                                                                                                                                                                                                                                98WO-US012209
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                                                                                                                                                                          RL;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         entry)
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77.0%;
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formulations - compris
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Pred. No. 4.2e-39;
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The present invention relates to a stable aqueous ph formulation, comprising an antibody not subjected to lyophilization, an acetate buffer of pH 4.8-5.5, a s polyol. The invention is useful for treating hemorrh injury, e.g. resulting from burns, stroke including
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The sequence is that of the heavy chain of a humanized murine anti-CD18 antibody rhuMab CD18. It can be used for the treament of disorders which include haemorrhagic shock, thermal injury (such as that resulting from burns) stroke (including ischaemic and haemorrhagic stroke) and myocardial infarction. The antibody formulation can be stabilised at a temperature of 2-8 deg. C for at least one year or at a temperature of deg. C for at least one month and is stable following freezing and deg. C for at least one month and is stable following freezing and
                                                                                                                                                                                                                                      Stable aqueous pharmaceutical formulation for treating hemorrhagic shock, thermal injury, stroke, and myocardial infarction, comprises an antibody
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                       A method has been developed for recovering a polypeptide comprising: (a) exposing a composition comprising a polypeptide to a reagent which binds to or modifies the polypeptide, where the reagent is immobilized on a solid phase; and (b) passing the composition through a filter bearing an opposite charge to the reagent so as to remove leached reagent from the composition. The present invention also describes a method for modifying a precursor antibody comprising a leucine zipper by exposing the precursor antibody to a protease immobilized on a solid support so that the contract of the process that the process that the contract of the process that the proce
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242. .277
/label= leucine_zip
      to a r
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77.0%;
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Pred. No. 4.4e-39;
7; Mismatches 15
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                                                                                                                                                                                                                                                                                                                                                                                                                            cell cultures
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port so that purify
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RESULT 9
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             proteins from cell cultures. They are especially useful for isolating antibodies. The methods overcome the problem of reagent leakage into the protein as is the case in prior art immobilized modification systems. Using a opposite charge filter the reagent can be excluded from the sample, preventing contamination. The present sequence represents the recombinant humanised anti-CD18 antibody (rhwMAb CD18) heavy chain, while used in an example from the present invention
                                                                                New stable aqueous antibody subjected to lyophilisation,
                                                                                                              WPI;
                                                                                                                              Weissburg
                                                                                                                                                                                                                                                                 Key
                                                                                                                                                                                                                                                                                  Synthetic
                                                                                                                                                                                                                                                                                                                                                   pS1130 expression
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence
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                                                                                                                                                                             13-JUN-1997;
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                                                                                                                                                                                                                                     WO9856418-A1
                                                                                                                                                                                                                                                      Peptide
                                                                                                                                                                                                                                                                                           Mus musculus.
                                                                                                                                                                                                                                                                                                              hemorrhagic;
                                                                                                                                                                                                                                                                                                                                 Monoclonal antibody; heavy chain; humanized; myocardial infarction;
                                                                                                                                                                                                                                                                                                                                                                       08-JUN-1999
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                                                                                                                                                                                                                                                                                                                         ourns; thermal
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g RP, Wong 1
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                                                                                                                                                                                                                                                                                                                                                                                                         standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                          TVSSAS
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                                                                                                                                                                                                                                                                                                               stroke;
                                                                                                                                                                                                                                                                                                        injury; injury; injury;
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                                                                                                                                                                             97US-00874897
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1. .23
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                                                                                                                                                           INC.
                                                                                                                                                                                                                                                                                                                                                   cassette encoded rhuMAb CD18 heavy chain
                                                                                                                                                                                                                                                                                                                                                                                                          protein;
                                                                                                                                RL;
                                                                                                                                                                                                                                                                                                                                                                      entry)
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77.0%;
                                                                                                                                       Ongpipattanakul
                                                                                                                                                                                                                                                                                                                         ischemic;
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                                                                               formulations, a buffer ma:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 499.5; Pred. No. 5.1e
                                                                                                                                                                                                                                                                                                                         shock; ischaemic;
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No. 5.1
                                                                                maintaining the
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                                                                                                                                        Shahrokh
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at 4.5 -
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The sequence is that of the pS1130 encoded heavy chain of a humanize murine anti-CD18 antibody rhuMAb CD18. It can be used for the treame disorders which include haemorrhagic shock, thermal injury (such as resulting from burns), stroke (including ischaemic and haemorrhagic stroke) and myocardial infarction. The antibody formulation can be

humanized

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Disclosure; Fig 21A; 87pp;

English

**surfactant** 

and a polyol.

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Best Local
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A method has been developed for recovering a polypeptide comprising: (a) exposing a composition comprising a polypeptide to a reagent which binds to or modifies the polypeptide, where the reagent is immobilized on a solid phase; and (b) passing the composition through a filter bearing an opposite charge to the reagent so as to remove leached reagent from the composition. The present invention also describes a method for modifying a precursor antibody comprising a leucine zipper by exposing the precursor antibody to a protease immobilized on a solid support so that
                                                                                                             Example;
                                                                                                                                 New method for recovering polypeptides from cell cultures - with reagent and filtering to remove reagent.
                                                                                                                                                                                                           Blank GS,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Recombinant
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          06-APR-1999
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                                                                                                                                                                                                                                                                                 12-JUN-1998;
                                                                                                                                                                                                                                                                                                          17-DEC-1998
                                                                                                                                                                                                                                                                                                                                                                                                     Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                              murine monoclonal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAW30635 standard; protein; 300
                                                                                                                                                                                                                                                                                                                               WO9856808-A1
                                                                                                                                                                                                                                                                                                                                                                                                                           Mus sp.
                                                                                                                                                                                                                                                                                                                                                                                                                                                    chromatography.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   freezing
                                                                                                                                                                                                                                   (GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                                                                                                  Peptide
                                                                                                                                                                                   1999-060267/05.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NORFMORFTISVOKSTSTAYMOMNSLRAEDTAVYYCARWRGLNYGFDVRYFDVWGQGTLV
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                                                                                                                                                                                                          Narindray
                                                                                                                                                                                                                                                                                                                                                                                                                                                              humanised anti-CD18 antibody; rhuMAb CD18; leucine zipper; clonal antibody; muMAb H52; protein recovery; filtration;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  humanised
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Conservative
                                                                                                            4; 43pp;
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RESULT 11
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Matches 97
                    The present invention relates to a stable aqueous pharmaceutical formulation, comprising an antibody not subjected to prior lyophilization, an acetate buffer of pH 4.8-5.5, a surfactant and polyol. The invention is useful for treating hemorrhagic shock, tinjury, e.g. resulting from burns, stroke including ischemic and hemorrhagic stroke, myocardial infarction, inflammatory disorders adult respiratory distress syndrome (ARDS), hypovolemic shock, ul
                                                                                                                                                       Stable aqueous pharmaceutical formulation for treating hemorrhagic thermal injury, stroke, and myocardial infarction, comprises an ant not subjected to prior lyophilization.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        the protease removes the zipper. The methods can be used to purify proteins from cell cultures. They are especially useful for isolating antibodies. The methods overcome the problem of reagent leakage into the protein as is the case in prior art immobilized modification systems. By using a opposite charge filter the reagent can be excluded from the sample, preventing contamination. The present sequence represents the recombinant humanised anti-CD18 antibody (rhuMAb CD18) heavy chain, which
                                                                                                                                                                                                                                                                                                                                                                                                                                             Antibody; lyophilization; hemorrhagic myocardial infarction; inflammation.
                                                                                                                                 Disclosure; Fig
                                                                                                                                                                                                               WPI; 2001-136863/14
                                                                                                                                                                                                                                                                                                                                                                09-JAN-2001.
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                                                                                                                                                                                                                                                                                                           13-JUN-1997;
                                                                                                                                                                                                                                                                                                                                     12-JUN-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                   Homo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Protein encoded
                                                                                                                                                                                                                                                                                 (GETH )
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                l Similarity
97; Conserv
                                                                                                                                                                                                                                        Oeswein JQ,
g RP, Wong
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   standard;
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             rheumatoid
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                                                                                                                                21; 56pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  by pS 1130 expression
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                protein;
             arthritis
                                                                                                                                                                                                                                          RL;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              entry.
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77.0%;
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                                                                                                                                 English
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Pred. No. 5.5e-39;
7; Mismatches 19
             and B-cell
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    B
                                                                                                                                                                                                                                                       B
                                                                                                                                                                                                                                                                                                                                                                                                                                                          shock; thermal injury;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     cassette
             lymphomas
                                                                                                                                                                                                                                                         Shahrokh
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                                                                                                                                                                                                                                                      Wang
                          disorders such as
shock, ulcerative
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                                                                                                                                                                         gic shock,
antibody
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Best Local S
Matches 97
                                                                                                                                                          Producing culturing and heavy
The present sequence is that of an anti-CD18 antibody heavy chain including a heat stable enterotoxin II (STII) secretion signal and a C-terminal leucine zipper domain. This recombinant heavy chain is produced by host cells transformed with claimed vector pxCD18-7T3 (see ABZ82071). The plasmid contains 2 translational units that temporally separate the transcription of the antibody light (see ABP72745) and heavy chains. Expression of the light chain is under the control of the photo promoter, while expression of the heavy chain is under the control of the inducible TacII promoter. In an example from the invention, Escherichia coli 59A7 was transformed with pxCD18-7T3. The light chain was expressed initially
                                                                                                                                     Claim
                                                                                                                                                                                                           N-PSDB;
                                                                                                                                                                                                                                              Andersen
                                                                                                                                                                                                                                                                                                                   26-AUG-2002; 2002WO-US027220
                                                                                                                                                                                                                                                                                                                                              06-MAR-2003
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                                                                                                                                                                                                                                                                                                                                                                                                        Protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Escherichia coli.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CD18; antibody; heavy chain;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Anti-CD18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            11-JUN-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ABP72746;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Chimeric.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Unidentified.
                                                                                                                                                                                                                                                                     (GETH ) GENENTECH
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                                                                                                                                  52; Fig 3; 73pp; English.
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                                                                                                                                                          an antibody for treating cancer or autoimmune diseases the host cell under suitable conditions so that the lic chain are expressed in a sequential fashion.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    antibody
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/note= "heat stable enterotoxin II (STII)
24. .300
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                                                                                                                                                                                                                                                                                                                                                                                                                                                     Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    heavy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           entry)
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                                                                                                                                                                                                                                                                                                                                                                                           "Heavy chain-leucine zipper"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    chain-leucine zipper
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Pred.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            $
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No. 5.5
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The H52 antibody ca is for treating for interrupted blood :

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f the invention. The damage to the brain

d H52 antibody. ion. The method ne brain due to

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ischaemic stroke,

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caused by obstruction of

a main

Disclosure;

Page

9; 41pp;

English.

Administering an anti-CD18 increasing cerebral blood :

flow

and

reducing size

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brain

infarct.

stroke

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antibody to treat focal ischaemic

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Best Local S
Matches 97
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   and secreted into the periplasmic space. Heavy chain production was then induced by addition of IPTG, and F(ab')2 assembly occurred almost immediately. A yield of 4.6 +/- 0.5 y/l assembled F(ab')2 was obtained, nearly 2-fold increase on the yield obtained using a single promoter system. This is an example of the process of the invention for productic of recombinant antibodies in a host cell system, with temporally separated expression of the light and heavy chains. Properly assembled, soluble and functional antibodies (or their fragments) can be produced in high yields for diagnostic or therapeutic applications, including treatment of cancer or autoimmune diseases
                                                                                                                                                              (GETH )
                                                                                                                                                                                                                                                                                                     Homo
                                                                                                                                                                                                                                                                                                                                               Humanised antibody; HuH52; heavy chain; focal ischaemic stroke; emboli brain damage; anti-CD18 antibody; cerebral blood flow; thromboembolism transient ischaemic attack; thrombolytic therapy; thrombosis; therapy;
                                                                                                                                        Bednar
                                                                                                                                                                                                23-JAN-1996;
                                                                                                                                                                                                                     11-JAN-1997;
                                                                                                                                                                                                                                           31-JUL-1997
                                                                                                                                                                                                                                                                                                                Synthetic
                                                                                                                                                                                                                                                                                                                                     systemic hypoperfusion; cardiac arrest.
                                                                                                                                                                                                                                                                                                                                                                                               Heavy chain of full length
                                                                                                                                                                                                                                                                                                                                                                                                                                            AAW34505;
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                                                                                                                                                              GENENTECH INC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                standard; protein;
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                                                                                                                                        Gross
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                                                                                                                                                                                                                                                                                                                                                                                                                     entry)
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77.0%;
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                                                                                                                                                               STATE AGRIC
                                                                                                                                        Thomas
                                                                                                                                                                                                                                                                                                                                                                                               IgG2 humanised H52 antibody.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   450
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Pred. No. 5.5e
7; Mismatches
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                                                                                                                                                               COLLEGE
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.5e-39;
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                                                                                                                                                                                                                                                                                                                                                               thromboembolism;
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                                                                                                                                                                                                                                                                                                                                                                          embolism;
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CC cerebral artery, comprises administering anti-CD18 antibody to increase CC cerebral blood flow and/or reduce the size of the area of brain that CC dies, without removing the arterial obstruction. CD18, is upregulated in CC patients after ischaemic stroke or transient ischaemic attack (stroke CC symptoms last less than 24 hours with complete recovery). Treatment with CC anti-CD18 antibodies can increase cerebral blood flow and/or reduce brain CC infarct size, improving clinical outcomes in focal ischaemic strokes in CC mammals, particularly in humans. The method provides an alternative to CC thrombolytic therapy for treating ischaemic strokes caused by CC thrombolytic therapy for treating ischaemic strokes caused by CC therapy has been unsuccessful, is contra-indicated e.g. because such CC drugs exacerbate bleeding, or is unsuitable because of the time delay CC between stroke onset and diagnosis. Thrombolytic agents (e.g. tissue CC plasminogen activator) may also be administered before, after or CC simultaneously with the anti-CD18 antibody. Alternatively, the method can be used to treat the third main type of ischaemic stroke, systemic CC hypoperfusion, e.g. resulting from cardiac arrest or drowning
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Best Local
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                                                                                                                                                                                                                                                                                                                                                       US5914112-A
                                                                                                                                                                                                                                                                                                                                                                                                                                                     IGG; immunoglobulin G; CH1 domain; human; anti-CD18; IgG1; IgG2; IgG3;
IgG4; Kappa-CL domain; lambda-CL domain; focal ischaemic stroke;
Cerebropyrotective; cerebral artery obstruction; blood flow; infarct;
CD18 extracellular domain; endothelium; CD11b/CD18 complex dissociation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAY08755;
Disclosure; Col 31-34; 25pp; English
                                        Anti-CD18 antibodies in stroke.
                                                                                 WPI; 1999-370483/31.
                                                                                                                                                                                                                               23-JAN-1996;
                                                                                                                                                                                                                                                                                                                22-JUN-1999
                                                                                                                                                                                                                                                                                                                                                                                           Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             huH52 heavy
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                                                                                                                         Bednar MM,
                                                                                                                                                                  VERMONT & STATE AGRIC COLLEGE.
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                                                                                                                                                                                                                                                                                                                                                                                                                                      heavy
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77.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           chain protein fragment.
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Pred. No. 8.6e-39;
7; Mismatches 15;
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Matches
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22-JAN-1997;
17-FEB-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          This invention describes a method for improving the clinical outcome in focal ischaemic stroke by administering novel anti-CD18 antibody which has cerebroprotective properties. The invention particularly describes a method of treating focal ischaemic stroke caused by the obstruction of a main cerebral artery which comprises administering an anti-CD18 antibody to increase the blood flow or reduce the infarct size, where: (1) the antibody binds to an extracellular domain of CD18 and inhibits or reduces the ability of the cell expressing CD18 to bind to endothelium, (2) the antibody binds CD18 with an affinity of less than 4 nm, or (3) the antibody dissociates CD11b/CD18 complex. This sequence represents the
                                                                                                                                                                                                                                                                                                                                                   Human; humanised H52 antibody; huH52; cerebral blood flow; focal ischaemic stroke; main cerebral artery, tissue plasminogen activator; anti-CD18 antibody; stroke; acute ischaemic stroke; thrombolytic therapy; Fabv1b salvage receptor binding epitope; thromboembolic st
                                                                                                                      Bednar MM,
                                                                                                                                                                                                                                                          27-JUN-2002.
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                                                                                                                                                                                                                                                                                                              Synthetic.
                                                                                                                                                                                                                                                                                                                           Homo sapiens.
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                                                                                                                                               (GETH ) GENENTECH INC.
                                                                                                                                                                                                                              20-DEC-2000; 2000US-00811384
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97; Conserv
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                                                                                                                    Gross CE,
                                                                                                                                                                                                                                                                                                                                                                                                                                   humanised H52 antibody (huH52), heavy chain.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Conservative
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97US-00788800.
99US-00251652.
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77.0%;
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                                                                                                                      Gross
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Increasing cerebral blood flow and/or reducing infarct size in focal ischemic stroke using anti-CD18 antibody and tissue plasminogen activator is useful to improve clinical outcome in acute ischemic stroke.

Disclosure; Page

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Search completed: December 29, 2004, 19:35:44 Job time : 125.575 secs
                                                                                                                                                                                                                                                                                                                           Query Match 78.8%;
Best Local Similarity 77.0%;
Matches 97; Conservative "
                                                                                                                                                                                                                                                                                                                                                                                                                                               The invention relates to a method of increasing cerebral blood flow and/ or reducing infarct size in focal ischaemic stroke caused by obstruction of a main cerebral artery in a human, comprising co-administering tissue plasminogen activator and anti-CD18 antibody about 3-5 hours after the stroke. The method is used to improve the clinical outcome in acute ischaemic stroke and to provide an alternative to thrombolytic therapy for treating thromboembolic stroke, particularly where thrombolytic therapy has been unsuccessful or is contra-indicated. The present sequence represents the full length humanised H52 antibody (huH52), heavy chain used in the method of the invention
                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 450 AA;
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                                                                                                                        114 TVSSAS 119
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Pred. No. 8.6e-39;
7; Mismatches 15;
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